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	14.2 15.3	226 14.2 15.3 227 14.2 15.3	225 14.2 15.3	224 14.2 15.3	223 14.2 15.3	221 14.2 13.3	220 14.2 15.3	219 14.2 15.3	14.2 15.3	14.2 15.3	216 14.2 15.3	215 14.2 15.3	14.2 15.3	14.2 15.3	14.2 15.3	211 14.2 15.3	210 14.2 15.3	209 14.2 15.3	14.2 15.3	14.2 15.3	14.4 15.5	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	205 14 4 15 5	204 14.4 15.5	203 14.4 15.5	202 14.4 LD.D	202 14 4 10 0	201 14.4 15.5	14.4 15.5	100 Hate at 150.0	199 14 4 15 5	14.4 15.5	19/ 14.4 LD.5	107 14 4 15 5	196 14.4 15.5	14.4 15.5	14.4 15.5	1 + · · · · · · · · · · · · · · · · · ·	193 14 4 15 5	192 14.4 15.5	191 14.4 L5.5	101	190 14 4 15 5	189 14.4 15.5	14.4 15.5	F# .# 10.0	1000	186 14 4 15 5	185 14.4 15.5	184 14.4 15.5	183 14.4 15.5	100 Hat. at 10.0	180 14 4 15 5	181 14.4 15.5	180 14.4 15.5	1/9 14.4 LO.5	110 H	178 144 155	14.4 L5.5	1 H	1/ / 10 0	14.4 15.5	1/4 14.4 15.5	111	173 1// 155	172 14.4 15.5	1/1 14.4 15.5	1/U 14.4 15.5	107 14.4 LO.D	160 14.4 10.0	168 144 155	14.4 15.5	100 14.4 15.5	100 Hat. at 1000	165 14 4 15 5	164 14.4 15.5	163 14.4 15.5	102 14.4 ID.D	160 14 4 10.0	161 14 4 15 5	160 14.4 15.5	159 14.4 15.5	14.4 15.5	14.4	157 14.4 15.5	156 14.6 15.7	
	14.2 15.3 38	226 14.2 15.3 35 227 14.2 15.3 37	225 14.2 15.3 34	224 14.2 15.3 33	223 14.2 15.3 33	221 14.2 13.3 32 222 14.2 15.3 32	220 L4.2 L5.3 3L	219 14.2 15.3 31	14.2 15.3 31	14.2 15.3 31	216 14.2 15.3 30	215 14.2 15.3 30	14.2 15.3 28	14.2 15.3 28	14.2 15.3 26	211 14.2 15.3 24	210 14.2 15.3 20	209 14.2 15.3 20	14.2 15.3 20	14.2 15.3 20	14.4 15.5 50	14.4 HU.U	205 14 4 15 5 50	204 14.4 15.5 50	203 14.4 15.5 50	202 14.4 LO.O 49	100 14 10 0 40	201 14.4 15.5 49	14.4 15.5 48	DDD 11-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1	199 14 4 15 5 48	14.4 15.5 48	19/ 14.4 LD.D 48	107 14 15 5	196 14.4 15.5 47	14.4 15.5 47	14.4 L5.5 47	14.# ±0.0 #/	193 14 4 15 5 47	192 14.4 15.5 46	191 14.4 15.5 46	101	190 14 4 15 5 44	189 14.4 15.5 43	14.4 15.5 41	14.4 LU.U 40	100	186 14 4 15 5 40	185 14.4 15.5 39	184 14.4 15.5 37	103 14.4 15.5 35	100 14.4 FU.U	182 14 4 15 5 34	181 14.4 15.5 34	180 14.4 15.5 32	1/9 14.4 15.5 31	110 14 14 14 14 14 14 14 14 14 14 14 14 14	178 144 155 31	14.4 15.5 31	11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	1/ / 18 8	14.4 15.5 29	1/4 14.4 15.5 29	1110 1110 100	172 1/ / 15 5 56	172 14.4 15.5 26	171 14.4 15.5 25	1/U 14.4 15.5 25	130 14.4 15.5 25	160 14 4 15 5	168 144 155 25	14.4 15.5 24	100 14.4 15.5 24	100 Hat. at 10.0 Na	165 14 4 15 5 24	164 14.4 15.5 24	163 14.4 15.5 24	102 14.4 15.5 24	160 14 4 10.0 60	161 14 4 15 5 20	160 14.4 15.5 20	159 14.4 15.5 20	14.4 15.5 19	14.4 10.0 17	157 14.4 15.5 17	156 14.6 15.7 50	
	14.2 15.3 38 18	226 14.2 15.3 35 20 227 14.2 15.3 37 22	225 14.2 15.3 34 22	224 14.2 15.3 33 24	223 14.2 15.3 33 24	221 14.2 15.3 32 24	220 14.2 15.3 31 22	219 14.2 15.3 31 20	14.2 15.3 31 20	14.2 15.3 31 20	216 14.2 15.3 30 22	215 14.2 15.3 30 20	14.2 15.3 28 21	14.2 15.3 28 21	14.2 15.3 26 24	211 14.2 15.3 24 19	210 14.2 15.3 20 22	209 14.2 15.3 20 22	14.2 15.3 20 20	14.2 15.3 20 19	14.4 15.5 50 22	11.1	205 14 4 15 5 50 22	204 14.4 15.5 50 22	203 14.4 15.5 50 22	202 II.4 ID.D 49 22	100 14 15 5 10 00 10 10 10 10 10 10 10 10 10 10 10	201 14.4 15.5 49 19	14.4 15.5 48 22	DDD 14.4 FU.U 40 61	199 14 4 15 5 48 21	14.4 15.5 48 19	197 14.4 LD.D 48 LB	107 14 15 5 40 10	196 14.4 15.5 47 21	14.4 15.5 47 19	14.4 L5.5 47 18	14.# LU.U #/ LU	193 14 4 15 5 47 15	192 14.4 15.5 46 24	191 14.4 L5.5 46 21	100 Hann HOLD AND HOLD HOLD HOLD HOLD HOLD HOLD HOLD HOL	190 14 4 15 5 44 16	189 14.4 15.5 43 16	14.4 15.5 41 22	T#.# LJ.J 40 LY	101 Hair HOLD 10 HA	186 14 4 15 5 40 14	185 14.4 15.5 39 18	184 14.4 15.5 37 24	103 14.4 15.5 35 ZI	102 14.4 10.0 04 10	180 14 4 15 5 34 19	181 14.4 15.5 34 18	180 14.4 15.5 32 15	1/9 14.4 LD.D 31 20	110 11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	178 144 155 31 10	14.4 15.5 31 14	11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	14 4 15 5 10 10	14.4 15.5 29 18	1/4 14.4 15.5 29 18	111	172 1// 155 26 20	172 14.4 15.5 26 20	1/1 14.4 15.5 25 24	1/U 14.4 15.5 25 24	102 14.4 13.3 23 24	160 14 4 10.0 60 64	168 14 4 15 5 25 24	14.4 15.5 24 24	100 14.4 L5.5 24 24	100 Hat. at 10.0 Nat Nat	165 14 4 15 5 24 24	164 14.4 15.5 24 24	163 14.4 15.5 24 24	102 14.4 15.5 24 22	160 14.4 10.0 20 24	161 14 4 15 5 20 24	160 14.4 15.5 20 20	159 14.4 15.5 20 14	14.4 15.5 19 19	14.4 HO.O HY HO	157 14.4 15.5 17 16	156 14.6 15.7 50 22	
	14.2 15.3 38 18	226 14.2 15.3 35 20 227 14.2 15.3 37 22	225 14.2 15.3 34 22	224 14.2 15.3 33 24	223 14.2 15.3 33 24	221 14.2 15.3 32 24	220 14.2 15.3 31 22	219 14.2 15.3 31 20	14.2 15.3 31 20	14.2 15.3 31 20	216 14.2 15.3 30 22	215 14.2 15.3 30 20	14.2 15.3 28 21	14.2 15.3 28 21	14.2 15.3 26 24	211 14.2 15.3 24 19	210 14.2 15.3 20 22	209 14.2 15.3 20 22	14.2 15.3 20 20	14.2 15.3 20 19	14.4 15.5 50 22	11.1	205 14 4 15 5 50 22	204 14.4 15.5 50 22	203 14.4 15.5 50 22	202 II.4 ID.D 49 22	100 14 15 5 10 00 10 10 10 10 10 10 10 10 10 10 10	201 14.4 15.5 49 19	14.4 15.5 48 22	DDD 14.4 FU.U 40 61	199 14 4 15 5 48 21	14.4 15.5 48 19	197 14.4 LD.D 48 LB	107 14 15 5 40 10	196 14.4 15.5 47 21	14.4 15.5 47 19	14.4 L5.5 47 18	14.# LU.U #/ LU	193 14 4 15 5 47 15	192 14.4 15.5 46 24	191 14.4 L5.5 46 21	100 Hann HOLD AND HOLD HOLD HOLD HOLD HOLD HOLD HOLD HOL	190 14 4 15 5 44 16	189 14.4 15.5 43 16	14.4 15.5 41 22	T#.# LJ.J 40 LY	101 Hair HOLD 10 HA	186 14 4 15 5 40 14	185 14.4 15.5 39 18	184 14.4 15.5 37 24	103 14.4 15.5 35 ZI	102 14.4 10.0 04 10	180 14 4 15 5 34 19	181 14.4 15.5 34 18	180 14.4 15.5 32 15	1/9 14.4 LD.D 31 20	110 11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	178 144 155 31 10	14.4 15.5 31 14	11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	14 4 15 5 10 10	14.4 15.5 29 18	1/4 14.4 15.5 29 18	111 11 11 10 10 10	172 1// 155 26 20	172 14.4 15.5 26 20	1/1 14.4 15.5 25 24	1/U 14.4 15.5 25 24	102 14.4 13.3 23 24	160 14 4 10.0 60 64	168 14 4 15 5 25 24	14.4 15.5 24 24	100 14.4 L5.5 24 24	100 Hat. at 10.0 Nat Nat	165 14 4 15 5 24 24	164 14.4 15.5 24 24	163 14.4 15.5 24 24	102 14.4 15.5 24 22	160 14.4 10.0 20 24	161 14 4 15 5 20 24	160 14.4 15.5 20 20	159 14.4 15.5 20 14	14.4 15.5 19 19	14.4 HO.O HY HO	157 14.4 15.5 17 16	156 14.6 15.7 50 22	
	14.2 15.3 38	226 14.2 15.3 35 20 227 14.2 15.3 37 22	225 14.2 15.3 34 22	224 14.2 15.3 33 24	223 14.2 15.3 33 24	221 14.2 15.3 32 24	220 14.2 15.3 31 22	219 14.2 15.3 31 20	14.2 15.3 31 20	14.2 15.3 31 20	216 14.2 15.3 30 22	215 14.2 15.3 30 20	14.2 15.3 28 21	14.2 15.3 28 21	14.2 15.3 26 24	211 14.2 15.3 24 19	210 14.2 15.3 20 22	209 14.2 15.3 20 22	14.2 15.3 20 20	14.2 15.3 20 19	14.4 15.5 50 22	11.1	205 14 4 15 5 50 22	204 14.4 15.5 50 22	203 14.4 15.5 50 22	202 II.4 ID.D 49 22	100 14 15 5 10 00 10 10 10 10 10 10 10 10 10 10 10	201 14.4 15.5 49 19	14.4 15.5 48 22	DDD 14.4 FU.U 40 61	199 14 4 15 5 48 21	14.4 15.5 48 19	197 14.4 LD.D 48 LB	107 14 15 5	196 14.4 15.5 47 21	14.4 15.5 47 19	14.4 L5.5 47 18	14.# LU.U #/ LU	193 14 4 15 5 47 15	192 14.4 15.5 46 24	191 14.4 L5.5 46 21	100 Hann HOLD AND HOLD HOLD HOLD HOLD HOLD HOLD HOLD HOL	190 14 4 15 5 44 16	189 14.4 15.5 43 16	14.4 15.5 41 22	T#.# LJ.J 40 LY	101 Hair HOLD 10 HA	186 14 4 15 5 40 14	185 14.4 15.5 39 18	184 14.4 15.5 37 24	103 14.4 15.5 35 ZI	102 14.4 LU.U U4 LO MAVUZU	180 14 4 15 5 34 19	181 14.4 15.5 34 18	180 14.4 15.5 32 15	1/9 14.4 LD.D 31 20	110 11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	178 144 155 31 10	14.4 15.5 31 14	11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	14 4 15 5 10 10	14.4 15.5 29 18	1/4 14.4 15.5 29 18	111 11 11 10 10 10	172 1// 155 26 20	172 14.4 15.5 26 20	1/1 14.4 15.5 25 24	1/U 14.4 15.5 25 24	102 14.4 13.3 23 24	160 14 4 10.0 60 64	168 14 4 15 5 25 24	14.4 15.5 24 24	100 14.4 L5.5 24 24	100 Hat. at 10.0 Nat Nat	165 14 4 15 5 24 24	164 14.4 15.5 24 24	163 14.4 15.5 24 24	102 14.4 15.5 24 22	160 14.4 10.0 20 24	161 14 4 15 5 20 24	160 14.4 15.5 20 20	159 14.4 15.5 20 14	14.4 15.5 19 19	14.4 HO.O HY HO	157 14.4 15.5 17	156 14.6 15.7 50 22	
	14.2 15.3 38 18	226 14.2 15.3 35 20 227 14.2 15.3 37 22	225 14.2 15.3 34 22	224 14.2 15.3 33 24	223 14.2 15.3 33 24	221 14.2 15.3 32 24	220 14.2 15.3 31 22	219 14.2 15.3 31 20	14.2 15.3 31 20	14.2 15.3 31 20	216 14.2 15.3 30 22	215 14.2 15.3 30 20	14.2 15.3 28 21	14.2 15.3 28 21	14.2 15.3 26 24	211 14.2 15.3 24 19	210 14.2 15.3 20 22	209 14.2 15.3 20 22	14.2 15.3 20 20	14.2 15.3 20 19	14.4 15.5 50 22	11.1	205 14 4 15 5 50 22	204 14.4 15.5 50 22	203 14.4 15.5 50 22	202 II.4 ID.D 49 22	100 14 15 5 10 00 10 10 10 10 10 10 10 10 10 10 10	201 14.4 15.5 49 19	14.4 15.5 48 22	DDD 14.4 FU.U 40 61	199 14 4 15 5 48 21	14.4 15.5 48 19	19/ 14.4 LD.D 48 LB	107 14 15 5 40 10	196 14.4 15.5 47 21	14.4 15.5 47 19	14.4 L5.5 47 18	14.# LU.U #/ LU	193 14 4 15 5 47 15	192 14.4 15.5 46 24	191 14.4 L5.5 46 21	100 Hann HOLD AND HOLD HOLD HOLD HOLD HOLD HOLD HOLD HOL	190 14 4 15 5 44 16	189 14.4 15.5 43 16	14.4 15.5 41 22	T#.# LJ.J 40 LY	101 Hair HOLD 10 HA	186 14 4 15 5 40 14	185 14.4 15.5 39 18	184 14.4 15.5 37 24	103 14.4 15.5 35 ZI	102 14.4 10.0 04 10	180 14 4 15 5 34 19	181 14.4 15.5 34 18	180 14.4 15.5 32 15	1/9 14.4 LD.D 31 20	110 11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	178 144 155 31 10	14.4 15.5 31 14	11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	14 4 15 5 10 10	14.4 15.5 29 18	1/4 14.4 15.5 29 18	111 11 11 10 10 10	172 1// 155 26 20	172 14.4 15.5 26 20	1/1 14.4 15.5 25 24	1/U 14.4 15.5 25 24	102 14.4 13.3 23 24	160 14 4 10.0 60 64	168 14 4 15 5 25 24	14.4 15.5 24 24	100 14.4 L5.5 24 24	100 Hat. at 10.0 Nat Nat	165 14 4 15 5 24 24	164 14.4 15.5 24 24	163 14.4 15.5 24 24	102 14.4 15.5 24 22	160 14.4 10.0 20 24	161 14 4 15 5 20 24	160 14.4 15.5 20 20	159 14.4 15.5 20 14	14.4 15.5 19 19	14.4 HO.O HY HO	157 14.4 15.5 17 16	156 14.6 15.7 50 22	
	14.2 15.3 38 18 AAT91908	226	225 14.2 15.3 34 22 AAH44703	224 14.2 15.3 33 24 ABK29277	223 14.2 15.3 33 24 ABR/11037	221 14.2 15.3 32 24 ABK1/322	220 14.2 15.3 31 22 AAI29566	219 14.2 15.3 31 20 AAX06542	14.2 15.3 31 20 AAX39358	14.2 15.3 31 20 AAX39205	216 14.2 15.3 30 22 AAS06509	215 14.2 15.3 30 20 AAX57295	14.2 15.3 28 21 AAZ52344	14.2 15.3 28 21 AAA37826	14.2 15.3 26 24 ABK69029	211 14.2 15.3 24 19 AAV65086	210 14.2 15.3 20 22 AAI68434	209 14.2 15.3 20 22 AAI68431	14.2 15.3 20 20 AAX93902	14.2 15.3 20 19 AAV64072	14.4 15.5 50 22 AAL33709	14 - 4 - 10 10 00 AMAD3/00	205 14 4 15 5 50 22 33123200	204 14.4 15.5 50 22 AAL32056	203 14.4 15.5 50 22 AAL32055	202 14.4 13.3 49 22 AAF 03322	000 14 4 15 5 10 00 35 15 15 15 15 15 15 15 15 15 15 15 15 15	201 14.4 15.5 49 19 AAV19177	14.4 15.5 48 22 AAS12833	14.4 13.3 40 Z1 AAA3/0/0	199 14 4 15 5 48 21 3357679	14.4 15.5 48 19 AAV47545	19/ 14.4 15.5 48 18 AAT/1/22	107 14 4 16 6 40 10 10 11 11 100 100 100 100 100 100	196 14.4 15.5 47 21 AAZ65831	14.4 15.5 47 19 AAX79243	14.4 15.5 47 18 AAT51661	14.4 13.3 4/ 13 AAQ/9244	193 14 15 5 47 15 33079344	192 14.4 15.5 46 24 AAS18520	191 14.4 15.5 46 21 AAA54317	The state of the s	190 14 4 15 5 44 16 33005160	189 14.4 15.5 43 16 AATO3423	14.4 15.5 41 22 AAI64806	TH. H LOUD 40 LY MAYOLI4/	100 First Ford to the PANCOSTIO	186 14 4 15 5 40 14 33053410	185 14.4 15.5 39 18 AATS1263	184 14.4 15.5 37 24 ABA93056	163 14.4 15.5 35 21 AAA10066	TO THE HOLD OF TO MAY UZ333	182 14 15 5 24 10 331703333	181 14.4 15.5 34 18 AATGGG16	180 14.4 15.5 32 15 AAQ77967	1/9 14.4 15.5 31 20 AAX06462	THE PARTY OF THE P	178 14 4 15 5 31 10 ANY 67761	14.4 15.5 31 14 AAQ44026	THE TOTAL OF ST PRINCIPLE	10 10 10 10 10 10 10 10 10 10 10 10 10 1	14.4 15.5 29 18 AATBQOB3	1/4 14.4 15.5 29 18 AAT89082	THE RESERVE TO SECURITION OF THE PROPERTY OF T	173 1/ / 18 5 26 20 3 3 3 2 5 2 2 2	172 14.4 15.5 26 20 AAY78418	1/1 14.4 15.5 25 24 ABN05074	1/0 14.4 15.5 25 24 ABN05073	102 14.4 13.5 25 24 ABQ12994	100 14 4 10 0 00 00 000 000 000 000 000	168 14 4 15 5 25 24 ABO 2053	14.4 15.5 24 24 ABQ11458	14.4 15.5 24 24 ABQ11417	100 14.4 10.0 24 24 ABQ00130	165 1/ / 15 5 3/ SECONDISC	164 14.4 15.5 24 24 ABOOKO89	163 14.4 15.5 24 24 ABO00576	102 14.4 15.5 24 22 AAH76860	160 14 A 15 5 24 20 34 ABRULDON	161 14 4 15 5 20 24 NOVELECO	160 14.4 15.5 20 20 AAZ21785	159 14.4 15.5 20 14 AAQ50576	14.4 15.5 19 19 AAV35420	1/ 15 5 10 10 PARCYCHOL	157 14.4 15.5 17 16 AAO95462	156 14.6 15.7 50 22 AAL33945	
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PCR primer for mou
             Mismatch probe spe
                                       Mouse gamma-actin
                                                                     Gamma-actin forwar
                                                                                                                           PCR primer UD2S160
Primer for 3' end
                                                                                                                                                                  Human MDMX antisen
                                                                                                                                                                                  Human
                                                                                                                                                                                                       Single nucleotide
                                                                                                                                                                                                                        Single nucleotide
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                                                       Reverse primer for
                                                                               Murine gamma actin
                                                                                              Reverse PCR primer
                                                                                                            A. thaliana NIM1 h
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                                                                                                                                                                                                                                                                                                         Human
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ALIGNMENTS

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RESULT 1
                                                                                                                       METH; metalloprotease; thrombospondin; anglogenesis inhibition; cancer therapy; benign tumour; ocular anglogenic disease; PCR primer; rheumatoid arthritis; psoriasis; wound healing; endometriosis; vasculogenesis; yranulation; hypertrophic scar; nonunion fracture; scleroderma, trachoma; vascular adhesion; myocardial angiogenesis; coronary collateral; cerebral collateral; arteriovenous malformation; ischaemic limb angiogenesis; osler-Webber syndrome; wound granulation; plaque neovascularisation; telangiectasia; haemophiliac joint; EST; angiotibroma; fibromuscular dysplasia; expressed sequence tag;
                                     W0200071577-A1.
30-NOV-2000
                                                                         Homo sapiens.
                                                                                                                Crohn's disease;
                                                                                                                                                                                                                                                                                                                             pFlag-CMV-5a:METH1 PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                19-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC90373 standard; DNA; 43 BP.
                                                                                                                                                                                                                                                                                                                                                                                                            AAC90373;
                                                                                                            atherosclerosis; birth control; ss.
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                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                               Query Match
                                       Bone morphogenic protein; bBMP-1; probes; cartilage formation; bone formation; osteogenic cpds; prodontal disease; ss.
                                                                                       Probe (32n) for the detection Bovine Bone Morphogenic Protein-1 sequence
                                                                                                                                                         AAN80618;
                                                                                                                         08-OCT-1990 (first entry)
                                                                                                                                                                                        AAN80618 standard; DNA; 32 BP
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 43 BP; 10 A; 11 C; 10 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                       neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, fibromuscular dysplasia, wound granulation, Crohn's disease or atherosclerosis. METH can also be used in birth control. METH can also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to human METH1 and METH2, (ME for metalloprotease and TH for thrombospondin; see ARB50002 and ARB50003). The present sequence is a PCR primer used in the present invention. METH can be used for inhibiting anglogenesis in an individual, and for treating cancer, benign tumours, an ocular anglogenic disease, rheumatoid arthritis, psoriasis, delayed wound healing, endometriosis,
                                                                                                                                                                                                                                                                                                                                                                                                                         be used in diagnostic methods for the prognosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vasculogenesis, granulations, hypertrophic scars, nonunion fractures, scleroderma, trachoma, vascular adhesions, myocardial angiogenesis, coronary collaterals, cerebral collaterals, arteriovenous malformations,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            METH1 and METH2 polynucleotides and encoded polypeptides, used to inhibit anglogensis in the treatment of disorders such as cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-025136/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Iruela-Arispe L, Hastıy:
Fornwald JA, Terrett JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ischaemic limb angiogenesis, Osler-Webber syndrome, plaque
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 52; Page 478; 768pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rheumatoid arthritis and psoriasis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-FEB-2000; 2000US-0183792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-AUG-1999;
22-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAY-2000; 2000WO-US14462.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (TERR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JONA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (IRUE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-)
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                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                  24;
                                                                                                                                                                                                                                                                   7 GGTACCACTGCATTCTGCCATTGTGCAAAAGTCT 40
                                                                                                                                                                                                                                                                                                6 GGGAGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TRUBEN S M.
JONAK Z L.
TRULLI S H.
FORNWALD J A.
                                                                                                                                                                                                                                                                                                                                                  Similarity
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BETH ISRAEL DEACONESS MEDICAL CENT.
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                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0147823
99US-0373658
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                                                                                                                                                                                                                                                                                                                                            19.48;
                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                            Score 18;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                              DB 22; Length 43;
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RESULT 3
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Best Local
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misc_binding
                                                                                                                                                                                                                                                        misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cancer cells detection oligonucleotide A18-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAL45727;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 32 BP; 7 A; 11 C; 9 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     See also AAN80619-N80636 and AAN81963-64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This probe is a relatively long (32 nucleotides) "guessmer" [See J.J. Toole et al, Nature, 312:342-347 (1984)]. The regions of the amino acid sequence used for probe design are chosen by avoiding highly degenerate codons where possible. The probe is radiolabelled with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; ; 7pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bone morphogenic proteins - obtd. using recombinant DNA and used for inducing cartilage and bone formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1988-021565/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENE-) GENETICS INST INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-MAR-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JUN-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JAN-1988
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                                                                                   protein_bind
                                                                                                                                                                    misc_binding
                                                                                                                                                                                                                                                                                                                                         misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumour cell detection; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL45727 standard; DNA; 49 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polynucleotide kinase and 32P-gamma-ATP and used to screen a bovine cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 CTGCCTGGGTGACAATGCCTTGGTT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     extranuclear DNA; stem-loop; DNA-binding protein; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTTCCTGGGCGACATTGCCCCTGGAT 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosen VA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           87US-0031346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                87WO-US01537.
/bound_moiety= "NKX2.5"
27..29
                                                                                   /bound_moiety= "nucleotides 29-27 of itself" 27..33
                                                 /*tag=
                                                                                                                                                                       /bound_moiety= "nucleotides 33-31 of itself"
21..23
                                                                                                                                                                                                                                                              /bound_moiety= "nucleotides 36-35 of itself"
17..19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                           /*tag=
                                                                                                                                                                                                                                                                                                                                                                     /bound_moiety= "nucleotide 38 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /bound_moiety= "nucleotides 44-40 of itself"
                                                                                                                                                                                                                               /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17; I
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 32;
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                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting tumor cells from presence of specific stem-loop DNA molecules outside the nucleus, useful for diagnosis and monitoring of tumors - \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    misc_binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                        in tissue sections, biopsies, body fluids etc, and can be used in the diagnosis and monitoring of cancer. The present sequence is an extranuclear DNA sequence capable of being detected using the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method of detecting tumour cells, by detecting extranuclear DNA consisting of a single-strand with a double-stranded stem-loop structure containing at least 2 binding sites for DNA binding proteins. The method can be used to detect cancer cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 21; Page 10; 27pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-SEP-2000; 2000DE-1046318.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DE10046318-A1
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 49 BP; 12 A; 8 C; 11 G; 18 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-331116/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Abken H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-SEP-2000; 2000DE-1046318
                                                                                Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                        Human silent SNP containing nucleic acid SEQ:2482.
                                                                                                                                                                     09-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-MAR-2002.
                           WO200140521-A2
                                                      Homo sapiens
                                                                                                                                                                                                 AAI75541;
                                                                                                                                                                                                                             AAI75541 standard; DNA; 50
                                                                                                                                                                                                                                                                                                                                                                                                                                              the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ABKE/) ABKEN H.
                                                                                                                                                                                                                                                                                                                                                                         ocal
                                                                                                                                                                                                                                                                                                                               5 TGGGAGCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTT 45
                                                                                                                                                                                                                                                                                                 2 TGGAAAAACTCCACTCACCTTGTTTCAAAGTGGGTTGGGTT 42
                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schinkoethe T;
                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= g
/bound_moiety= "nucleotides 23-21 of itself"
31..33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              40..44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /bound_moiety= "nucleotides 10-6 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /bound_moiety= "nucleotide 12 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /bound_moiety= "nucleotides 19-17 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bound_moiety= "nucleotides 15-14 of itself"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                                                         63.48;
                                                                                                                                                                                                                                                                                                                                                                                       18.3%;
                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                       Score 17;
                                                                                                                                                                                                                                                                                                                                                                         Pred. No.
                                                                                                                                                                                                                                                                                                                                                                         9.3e+03
                                                                                                                                                                                                                                                                                                                                                                                       DB 24; Length 49;
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07-JUN-2001

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(MIAC ) CANADA MIN FISHERIES & OCEANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative
                                     17-JUN-1994;
                                                                                                                                               GB2327679-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy. The polypeptides encoded by (I) may be used as antigens in t production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
                                                                        16-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequences (1), which contain single nucleotide polymorphisms (SNPs).

AAM5314 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (1) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases
                                                                                                              03-FEB-1999
                                                                                                                                                                                   Oncorhynchus nerka
                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                    histone H3; promoter; region; salmon; PCR primer;
                                                                                                                                                                                                                                                                         Oncorhynchus nerka GH5 PCR primer.
                                                                                                                                                                                                                                                                                                               08-JUN-1999
                                                                                                                                                                                                                                                                                                                                                  AAX07403;
                                                                                                                                                                                                                                                                                                                                                                                      AAX07403 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 50 BP; 13 A; 13 C; 13 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 811; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-NOV-2000; 2000US-0726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-2000; 2000WO-US32758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      37 CCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCAGA 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 CCATGGCGTGCACACAAGGGGACTGTCAATCACAGGCTTTCATA 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Leach M;
                                     94CA-2126138
                                                                        95GB-0012302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18.1%; Score 16.8; 61.4%; Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ..1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 50;
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polypeptide comprising a Us1.5 phosphorylation site, a candidate inhibitor and a kinase and compares the phosphorylation state of the polypeptide with that of a similar polypeptide contacted with the kinase in the absence of the candidate inhibitor. The inhibitor of Us1.5 phosphorylation is used to inhibit herpesvirus infection in a cell before or after the cell is contacted with herpesvirus. In particular the inhibitor is used to prevent herpes simplex virus infections. The present sequence represents a PCR primer used in
                                                                                                                                               substances able to inhibit herpesvirus infections.
                                                                                                                                                                                                                                                     Assay for detecting inhibitors of herpesvirus infections monitors the phosphorylation state of polypeptides with a Usl.5 phosphorylation site
                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequence is that of a salmon promoter PCR primer which was used in a process for the production of transgenic fish. This is used to increase the growth of fish in fish farming and can lead to more significant levels of gene expression in fish.
                                                                                                                                                            The present invention describes a screening assay to determine
                                                                                                                                                                                                   Example 1; Page 89; 143pp; English.
                                                                                                                                                                                                                                                                                                             WPI; 2000-619232/59.
                                                                                                                                                                                                                                                                                                                                                                                   (ARCH-) ARCH DEV CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-APR-2000; 2000WO-US09215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200061794-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Herpesvirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Herpesvirus; post translational modification; infection; inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Herpesvirus expression vector construction PCR primer P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC64543 standard; DNA; 33 BP
                                                                                                                                                                                                                                                                                                                                              Roizman B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-OCT-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 30 BP; 8 A; 8 C; 8 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Preparation of transgenic fish by transforming fish cell with expression vector - comprises combining homologous fish histone H3 or metallothionein gene promoter DNA sequence, growth hormone gene sequence and fish gene terminator DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC64543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Devlin RH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 10; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-084295/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 ATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 ATCCAGCCTGGATGACAATGACT 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 identification; Us1.5; phosphorylation site; virucide;
                                                                                                                                                                                                                                                                                                                                              Ogle WO;
                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0128500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.6; DB 20;
Pred. No. 1.1e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                               The assay uses a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                   Matches
                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 33 BP; 8 A; 9 C; 11 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the construction of herpesvirus expression vectors.
                                                                                                                                                                                                                                             AATO8176-T08219 represent amplification primers used in the construction of modified ligands. The modified ligands comprise two copies of a sequence joined to a linker. The sequences used in the ligand include interleukin-4 (IL-4), IL-2, granulocyte-colony stimulating factor (G-CSF) and granulocyte macrophage-colony stimulating factor (GM-CSF). The linker regions used are represented by AAR87726 and AAR87730. This sequence was used in conjunction with AATO8200 to amplify codons 39-127 of IL-2 for use in a modified ligand. These modified ligands are known as circularly permuted ligands. The modified ligands can be joined to
                                                                                                                                                     another protein (preferrably a cytotoxin or an antibody) using a spacer (see ARR87732-R87734). The circularly permuted ligands have greater specificity and binding affinity than the original ligand. They can be used in chimeric molecules to inhibit the growth of tumour cells in vivo, where the two copies of the protein sequence are joined to cytotoxins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Linker; interleukin-4; interleukin-2; IL-4; IL-2; G-CSF; GM-CSF; granulocyte-colony stimulating factor; circularly permuted ligand; cancer; therapy; granulocyte macrophage-colony stimulating factor;
                                                                                                                       Sequence 39 BP; 7 A; 10 C; 12 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                Example 5; Page 35; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interleukin(s) or colony-stimulating factors - with inc
binding specificity and affinity for inhibiting tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Circularly permuted ligands and chimeric mols. are modified interleukin(s) or colony-stimulating factors - with increased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-366354/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kreitman RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-0CT-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9527732-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymerase chain reaction; PCR; primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amplification primer BK138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-MAY-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT08201 standard; DNA; 39 BP.
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                                                                    FOCGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 AATGTCGGCCGTCCAGAAAACGT 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            64 ACTGTCGGGCCTCCAGAAAACAT 86
               43 TTTCAACACTATCGGAATGTGACTGTCGGGC 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                    Similarity
Conservative
                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pastan I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94US-0225224
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                                                                  17.8%;
71.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16.6;
                                                                                    Score 16.6;
                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                     Mismatches
                                                                      No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 21;
                                                                      .2e+04
                                                                                        DB 16;
                                                       9,
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                                                       Indels
                                                                                      Length 39;
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                                                       Gaps
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AAC90375
Вb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer therapy; benign tumour; ocular angiogenic disease; PCR primer; rheumatoid arthritis; psoriasis; wound healing; endometriosis; vasculogenesis; granulation; hypertrophic scar; nonunion fracture; vasculogenesis;
                  The present invention relates to human METH1 and METH2, (ME for metalloprotesse and TH for thrombospondin; see AB50002 and AB50003. The present sequence is a PCR primer used in the present invention. METH can be used for inhibiting angiogenesis in an individual, and for treating cancer, benign tumours, an ocular angiogenic disease, rheumatoid arthritis, psoriasis, delayed wound healing, endometriosis, vasculogenesis, granulations, hypertrophic scars, nonunion fractures, scleroderma, trachoma, vascular adhesions, myocardial angiogenesis, scleroderma, trachoma, vascular adhesions, myocardial angiogenesis, coronary collaterals, cerebral collaterals, arteriovenous malformations, ischaemic limb angiogenesis, Osler-Webber syndrome, plaque neovascularisation, telangiectasia, haemophiliae, joints, angiofibroma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Crohn's disease; atherosclerosis; birth control; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pC4Fc:METH1 PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC90375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC90375 standard; DNA; 43 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAY-2000; 2000WO-US14462.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          METH; metalloprotease; thrombospondin; angiogenesis inhibition;
                                                                                                                                                                                                                                                                                                  METH1 and METH2 polynucleotides and encoded polypeptides, used to inhibit angiogensis in the treatment of disorders such as cancer,
                                                                                                                                                                                                                                                                                                                                                                                  Iruela-Arispe L, Hasting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200071577-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-JUL-1999;
                                                                                                                                                                                                                                        Example 52; Page 479; 768pp; English.
                                                                                                                                                                                                                                                                          inhibit angiogensis in the treatment of disorders rheumatoid arthritis and psoriasis -
                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-025136/03
                                                                                                                                                                                                                                                                                                                                                                                                                                              (TERR/) TERRETT J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ( HUMA - )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l3-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IRUE/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JONA/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    35 TCTCAACACTCACCGGAGGTAACGGTGGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TRULLI S H.. FORNWALD J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RUBEN S M.
JONAK Z L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BETH ISRAEL DEACONESS MEDICAL CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HASTINGS G A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IRUELA-ARISPE L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0183792.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0171503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0318208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0147823.
                                                                                                                                                                                                                                                                                                                                                                                                        Hastings GA, Ruben SM, Jonak ZL, Trulli SH;
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fibromuscular dysplasia, wound granulation, Crohn's disease

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RESULT 9
AAZ66265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ន្តន្តន
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                    invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side
                                               N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the research in the sequence of th
                                                                                                                  effects from pharmaceutical agents acting on a disease as well as other
                                                                                                                                                                                                                                                                                                 AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the
                                  from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 361; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cohen D,
                                                                                                      treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis; single nucleotide polymorphism; SNP; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human map-related biallelic marker SEQ ID NO:612.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ66265;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ66265 standard; DNA; 47 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 43 BP; 11 A; 10 C; 9 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             atherosclerosis. METH can also be used in birth control. METH can also be used in diagnostic methods for the prognosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
es 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10 AGAACTGCATTCTGCCATTGTGCAAAAGTCT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9 AGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Blumenfeld M, Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-0109732.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0082614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-IB00822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 replace(24,C)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17.8%;
71.0%;

    Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.6;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
RESULT 11
AAQ98309
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                X P X P X X F T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Дb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ63029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SO
                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                       the procedure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1994-135610/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 47 BP; 12 A; 10 C; 14 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NEXA-) NEXAGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9408050-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ63029;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-SEP-1992;
                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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Three oligonucleotides (AAQ63028-Q63030) were used together in a variable template SELEX procedure (Experiment B) to produce ligands for basic fibroblast growth factor (bFGF). This is a primer used in
                                                                                                                                                                           Sequence 50 BP; 13 A; 13 C; 13 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                     Producing target specific nucleic acid ligands - by selection for high affinity then structure determination, esp directed against HIV proteins, thrombin or basic fibroblast growth factor
                                                                                                                                                                                                                                                                                                                                 Example 5; Page 106; 208pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gold LM, Janjic N, Tasset D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-NOV-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rev protein; tat protein; basic fibroblast growth factor; bFGF; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer used in SELEX procedure for production of bFGF ligands.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thrombin; coagulation; adhesion; blood clot; clot formation; ligand; binding; SELEX; human immunodeficiency virus; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ63029 standard; DNA; 50 BP
20 CACTATAGGGAGATGCCTGTCGAGCATGCTG 50
                                      49 CACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GAGCTGGGAGCACTCCATTCTGC 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 GGGCTGGGACCACAACATTCTGC 30
                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93US-0061691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93WO-US09296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-0973333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9205-0953694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers 11..26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= T7 promoter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                          17.8%;
71.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17.8%;
                                                                                                                               Score 16.6; DB 15;

 Mismatches

                                                                                                            Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 16.6;
Pred. No. 1.
                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tuerk C;
                                                                                                          No.
                                                                                                            1.3e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.3e+04;
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                                                                                         9;
                                                                                                                               Length 50;
                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                    Gaps
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                                                                                                                                                                                     CC synthesized. The random region was generated by utilising an equimolar CC mixture of the four nucleotides during oligonucleotide synthesis. The CC constant regions were designed to contain PCR primer annealing sites, CC allowing cDNA synthesis and containing a T7 RNA promoter region. An CC initial pool of RNA molecules was prepared by in vitro transcription of CC the double stranded DNA template. Transcription mixtures were incubated CC at 37 deg. C for 2-3 hours which resulted in a typical amplification of CC incubating with bFGF for 10 mins a buffer solution at 37 deg. C then CC incubating with bFGF for 10 mins a buffer solution at 37 deg. C then CC counds of selection, no additional improvement in binding was seen.

CC The experiment was repeated using the sequences given in AAQ98305-08, CC without heparin as a competitor for binding of randomised RNA to bFGF. CC The isolated ligands fell into two families, family 1 having a consensus CC sequence of CUAACCAGG and family two having the consensus sequence given
밁
                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequences given in AAQ98309-10 are primers which were used in the isolation of ligands to thrombin. The ligands were isolated using systematic evolution of ligands by exponential enrichment (SELEX). DNA templates such as the one in AAQ98308, containing a region of 30 random nucleotides flanked by constant sequence regions, were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SELEX primer 1 molecule for isolation of bFGF ligands (experiment B)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ98309 standard; DNA; 50 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NEXS-) NEXSTAR PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9521853-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       systematic evolution of ligands by exponential enrichment; SELEX; heparin; selection; region of homology; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Family 1; family 2; ligand; thrombin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ98309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identification of ligands to basic fibroblast growth factor and thrombin - which can be modified for increased in vivo stability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1995-293073/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-JUN-1990;
10-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-MAR-1994;
                                                                                                                                        Sequence 50 BP; 13 A; 13 C; 13 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 69; 236pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-FEB-1994;
                                                                                                                                                                             in AAQ98434.
                                                                                    Match 17.8%; Local Similarity 71.0%;
20
                                    49 CACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Janjic N, Tasset D;
CACTATAGGGAGATGCCTGTCGAGCATGCTG 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90US-0536428.
91US-0714131.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95WO-US01458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93US-0061691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94US-0195005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-0219012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers 11..26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/note= "T7 promoter"
                                                               0;
                                                                                      Score 16.6; DB 16; Pred. No. 1.3e+04;
                                                                       Mismatches
                                                                                                      Length 50;
                                                                       Indels
                                                                       0,
                                                                     Gaps
                                                                         0
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RESULT 12
AAL29115
                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antinicrobial; gene therap; vaccine; amylase; cancer; amyloid protein; anglopoisetin; apptosis related protein; conference; complement related protein; cytochrome; kinase; colony stimulating factor; complement related protein; cytochrome; kinase; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAL29115 standard; DNA; 50 BP.
                                                                                                                                                                                                                                    by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, anglopoletin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, grotein coupled receptors and thioesterases. The present sequence is one such oligonucleotide, The oligonucleotides and the peptides encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human SNP oligonucleotide #2323.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL29115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-465210/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-DEC-2000; 2000WO-US35498
                                                                                                                                   Sequence 50 BP; 4 A; 10 C; 25 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 2047; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shimkets RA, Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUL-2001.
                              Loca1
                                  Similarity
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0173419
                       17.8%;
64.1%;
0
                                  Score 16.6; DB 23
Pred. No. 1.3e+04
   Mismatches
                                                                     DB 22;
   14; Indels
                                                                     Length 50;
       0;
       Gaps
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B ρy

5 TGGGAGCACTCCATTCTGCCTGGGTGACAATGCCTTGGT 43

RESULT 13

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                                                                                                                                                                                                                                                   AAV29852/c
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                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                      5' VH primer for expression with linker peptide.
                           Synthetic
                                                     genetic immunisation; systemic lupus erythematosus; antibody;
autoimmune disease; tumour; cancer; H221; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a purified and isolated non-naturally occurring DNA ligands to basic fibroblast growth factor (bFGF). The ligands are useful as part of gene therapy treatments and for diagnosing pathogenesis of vascular diseases including initiation and progression of atherosclerosis, acute coronary syndromes, vein graft disease and restenosis following coronary angioplasty. The ligands have improved stability in vivo.
                                                                                      Intracellular targeting sequence; DNA vaccine;
                                                                                                                                                            28-SEP-1998 (first entry)
                                                                                                                                                                                                                               AAV29852 standard; DNA; 47 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 50 BP; 13 A; 13 C; 13 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel nucleic acid ligands to basic fibroblast growth factor that are useful as inhibitors of basic fibroblast growth factors and 2'-amino modified RNA ligands, exhibit increased in vivo stability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Janjic N, Gold L, Tasset D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 19; Column 36; 153pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-158583/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-JUN-1990;
10-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NEXS-) NEXSTAR PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US6177557-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               atherosclerosis; angioplasty; stability; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ligand; basic fibroblast growth factor; bFGF; gene therapy; vascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SELEX experiment oligonucleotide #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-APR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF70543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF70543 standard; DNA; 50 BP
                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                               49 CACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                20 CACTATAGGGAGATGCCTGTCGAGCATGCTG 50
                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94US-0219012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-0195005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9208-0973333.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91US-0714131.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90US-0536428.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96US-0687421
                                                                                                                                                                                                                                                                                                                                                                                                   17.8%;
71.0%;
                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                   Score 16.6; DB 22;
Pred. No. 1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             .3e+04;
                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                Length 50;
                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                      AC XXX DT XXX AC XXX XXX XXX XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           γQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
28-OCT-1999.
                                 WO9954500-A2
                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                  diagnosis; single nucleotide polymorphism; SNP; ds
                                                                                                                                                                                               genomic map: haplotype; phenotype; polymorphic base; genotyping;
haplotyping; hybridisation; identification; characterisation;
                                                                                                                                                                                                                               Human genome; biallelic marker; high density disequilibrium map;
                                                                                                                                                                                                                                                                  Human map-related biallelic marker SEQ ID NO:1452.
                                                                                                                                                                                                                                                                                                       10-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                       AAZ67105 standard; DNA; 47 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 47 BP; 11 A; 12 C; 11 G; 10 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in DNA vaccination studies. The invention provides a novel plasmid comprising a sequence that encodes an immunogenic target protein that includes, or is linked to, an intracellular targeting sequence (see AAW56512-14) that directs localisation of the target protein to e.g. the lysosome or ER. The plasmid is used as a DNA vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encode the VH or Fv region of H221. This antibody binds to murine renal mesangial cells and aortic endothelial cells, properties associated with pathogenic systemic lupus erythematosus antibodies. H221 VH and VL regions have been amplified by PCR (see AAV29841-54) and used to generate H221 VH and Fv coding sequences having signals for intracellular or extracellular expression. These were cloned into a genetic immunisation vector and administered to mice for use in DNA vaccination studies. The invention provides a novel plasmid
                                                                                                     variation
                                                                                                                                                                                                                                                                                                                                            AAZ67105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This 5' primer was used for the PCR amplification of DNA encoding the VH region of murine anti-DNA IL/LM (H221) antibody for expression with linker peptide. DNA constructs are provided that encode the VH or Fv region of H221. This antibody binds to murine encode the VH or Fv region of H221.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmid encoding immunogenic target protein - used in, e.g. protective or therapeutic vaccines against allergy, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 52; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    microbial infection or autoimmune disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-261198/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Madaio M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-APR-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9817323-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYPE-) UNIV PENNSYLVANIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Loca 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        55 CGGAATGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CAGRYTGYTGCAGCTGGACCTCCTGATAAGATTTAGAT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Weiner DB, Williams WV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96US-0029592
                                                                                               replace(24,T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97WO-US19545.
                                                                                                                  Location/Qualifiers
                                                            /standard_name= "single nucleotide polymorphism"
                                                                               /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17.6%;
60.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16.4; DB 19;
Pred. No. 1.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                    PF PFD PACE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side
                                                                                                                                                                                              Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antinicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; anglopoiettin; apptoosis related protein; catherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinase; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cohen D, Blumenfeld M, Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 47 BP; 16 A; 6 C; 16 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 540; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEST ) GENSET.
28-DEC-1999;
                                                                      05-JUL-2001.
                                                                                                           WO200147944-A2
                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                   Human SNP oligonucleotide #7388.
                                                                                                                                                                                                                                                                                                                                                                                       24-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                              AAL34180;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL34180 standard; DNA; 50 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        effects from pharmaceutical agents acting on a disease as well as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       map of the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel biallelic markers used to construct a high density disequilibrium
                                  28-DEC-2000; 2000WO-US35498.
                                                                                                                                                                               nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           36 CTGCTTCCCTACGTATGCCTTGGTTTCCATCCTA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 CTGCCTGGGTGACAATGCCTTGGTTTCAACACTA 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0082614
98US-0109732
99US-0173419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-IB00822.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17.6%;
67.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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RESULT 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 approtosis related proteins, cadherin, cyclin, polymerase, onogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and drave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic propositions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 3513; 4143pp; English.
                                                                                                                                                                                                                                                                                                           Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoiet:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-465210/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shimkets RA, Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                     09-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI75539;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 50 BP; 10 A; 17 C; 20 G; 3 T; 0 other;
               WPI; 2001-356160/37.
                                                                                                              30-NOV-1999; 99US-0168138.
29-NOV-2000; 2000US-0726173.
                                                                                                                                                                     30-NOV-2000; 2000WO-US32758
                                                                                                                                                                                                         07-JUN-2001.
                                                                                                                                                                                                                                          WO200140521-A2.
                                                                                                                                                                                                                                                                                                                                                                                 Human silent SNP containing nucleic acid SEQ:2480.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAI75539 standard; DNA; 50 BP
                                               Shimkets RA,
                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     госат
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17 CGGAGCGGCCAGGCCGGGCCTCCCCAAAAGCCTG 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  55 CGGAATGTGACTGTCGGGCCTCCAGAAAACATTG 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                   Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17.6%;
67.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.6e+04
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                                     Preparation of cDNA from RNA molecules - by annealing cDNA synthesis
switching oligo:nucleotide
                  primer to RNA, synthesising DNA and contacting with novel template
                                                                                                 WPI; 1997-363690/33.
                                                                                                                                            Chenchik A, Diatchenko L, Siebert P,
                                                                                                                                                                                                                                                   03-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                  10-JUL-1997.
                                                                                                                                                                                                                                                                                                                                                                                                  W09724455-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      misc_RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Template switching oligonucleotide; RNA/DNA hybrid; DNA preparation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Template switching oligonucleotide NalSuplM.
                                                                                                                                                                                                 (CLON-) CLONTECH LAB INC.
                                                                                                                                                                                                                                                                                                03-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA synthesis primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-MAR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT98491;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT98491 standard; DNA; 40 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 50 BP; 13 A; 15 C; 13 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           presence of polymorphic polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 810; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37 CCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCA 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9 CCATGGCGTGCACACAAGGGGACTGTCAATCACAGGCTTTCA 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         h 17.6%;
Similarity 61.9%;
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                                                                                                                                                                                                                                                   96US-0582562
                                                                                                                                                                                                                                                                                             97WO-US00368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers 38..40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA cloning; cDNA library construction; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. 1.6e+04
                                                                                                                                              Zhu Y;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC AAT98426-T98496 represent template switching oligonucleotides (TSO) used CC in the method of the invention is for CI in the method of the invention is for CC preparing DNA complementary to the 5'-end of an RNA molecule. The method CC comprises annealing a CDNA synthesis primer (such as AAT98497) to the RNA CM molecule and synthesising a first DNA strand complementary to at least a CM portion of the RNA molecule, and contacting the RNA molecule with a TSO CM aving a pre-selected nucleotide sequence at its 5'-end and at least 1 CM aving a pre-selected nucleotide sequence at its 5'-end and at least 1 CM aving a pre-selected nucleotide sequence at its 5'-end and at least 1 CM aving a pre-selected nucleotide sequence at its 5'-end of the 3'-end of the CC riboguanine residue at its 3'-end, where the TSO binds the 5'-end of the CC completie and serves as a todmitter for the extension of the 3'-end of CC cloning of full length cDNA, or fragments, that correspond to the CC complete sequence of the 5'-end of the mRNA molecule. It can be used to CC construct cDNA libraries from nanogram quantities of total or poly A+CC complementary to the 5'-end of the template RNA, while allowing full length on A practive selection against cDNA that are not complementary to the 5'-end of the template RNA, while allowing full length on A practive selection against cDNA that are not complementary to the S'-end of the template RNA, while allowing full length on A practive selection against cDNA that are not complementary to the S'-end of the template RNA, while allowing full length on A practive selection against cDNA that are not complementary to the S'-end of the mRNA while allowing full length on A practive selection against cDNA that are not complementary to the S'-end of the mRNA while allowing full length on A practical selection against cDNA that are not length on A practical selection against cDNA that are not length on A practical selection against cDNA that are not length of the selection against cDNA that are not length of th
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The invention relates to a template switching oligonucleotide (1) with pre-selected arbitrary nucleotide sequence at its 5' end, at least 3 ribonucleotide units at its 3' end with at least one nucleotide that forms a basepair(s) with a nucleotide(s) at least one nucleotide that strand of an RNA-CDNA intermediate and is capable of serving as a template for extension of the 3' end of the first cDNA strand. The oligonucleotides ARZ28627-28740 represent examples of oligonucleotides used in the method of the invention. The oligonucleotide is used for the synthesis of full length cDNA corresponding to the complete sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 12; Column 29-30; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Template switching oligonucleotide useful for the synthesis of full
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-589715/50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CLON-) CLONTECH LAB INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 9; Page 30; 39pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 GCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24; Conservative
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Similarity 64.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Siebert P, Chenchik A, Zhu Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96US-0582562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0778494.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. 1.7e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV76941/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 20
                               CC that protein encoding or regulatory regions of commercial, therapeutic or cc industrial importance can be obtained. Specifically, sequences which are clikely to encode antigens have been identified and these polypeptides can cc be used in a vaccine composition against S. aureus infection. The polypeptides can also be used in a kit for the immunodetection of cc saureus in a sample. S. aureus is implicated in numerous human diseases, cincluding cellulitis, eyelid infections, food poisoning, osteomyelitis, cc skin and surgical wound infections, scalded skin syndrome, etc. Organisms transformed with the DNA sequences can be used cfor recombinant production of the polypeptides. The new DNA sequences cc (and their fragments) are useful as primers or probes for isolating cc computer readable medium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 40 BP; 12 A; 9 C; 16 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5'-ends of mRNA molecules. It allows the production of cDNA libraries from nanogram quantities of total or polyA+ RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Staphylococcus aureus contig SEQ ID #2630.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-MAR-1999 (first entry)
                                                                                                                                                                                                                                                                                                        This sequence represents one of 5191 Staphylococcus aureus DNA sequences of the invention. The DNA sequences are recorded on a computer readable medium, preferably selected from a floppy or hard disk, random access memory (RAM), read-only memory (ROM) or CD-ROM. Homology searches using
                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 2293; 3271pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              polynucleotide(s) and proteins derived from Staphylococcus aureus
stored on computer readable medium and used in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-374922/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Barash SC, Choi GH, Dillon PJ, Fannon MR, Kunsch CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JUL-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP786519-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         toxic shock syndrome; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Computer readable medium; vaccine; S.aureus infection; immunodetection; cellulitis; eyelid infection; food poisoning; osteomyelitis; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV76941 standard; DNA; 50 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                               anti-S.aureus vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          skin infection; surgical wound infection; scalded skin syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        37 GCTGGCTGCCCACTCTGCGTTGTTACCACTGCTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 GCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 17.4%; Score 16.2;
Similarity 64.9%; Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-0009861.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97EP-0100117.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Sequence 50 BP; 16 A; 10 C; 14 G; 9 T; 1 other;

Matches 24;

Conservative

0; Mismatches

Indels Length 50;

0;

Gaps

0;

Score 16.2; DB 22 Pred. No. 1.9e+04;

DB 22;

Match 17.4%; Local Similarity 64.9%;

Sequence 50 BP; 20 A; 7 C; 5 G; 18 T; 0 other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         complement related protein; cytochrome; kinesin; cytokine; interfero interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL27838;
                                                                                                                                                                                                          proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded
                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-JUL-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL27838 standard; DNA; 50 BP
                                                                          by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney.
                                                                                                                                                                                                                                                                apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                       The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases amyloid proteins, anglopoletin,
                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 1679; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-465210/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-DEC-1999; 99US-0173419.
27-DEC-2000; 2000US-0173419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                   cancer, autoimmune diseases and infections .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CURA-) CURAGEN CORP.
                                         organisms.
                                                      leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        47 CTTTGGTTTAAACCCAATCGGCCTTTAAC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         37 CCTTGGTTTCAACACTATCGGAATGTGAC 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17.48;
72.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16.2; DB 18
Pred. No. 1.9e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 50;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interferon;
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CC The present invention relates to oligonucleotides encoding polymorphic CC variants of proteins related to amylases, amyloid proteins, angiopoietin, CC apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, CC histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, CC G-protein coupled receptors and thioesterases. The present sequence is CC one such oligonucleotide. The oligonucleotides and the peptides encoded CC by them may be used in the prevention, diagnosis and treatment of Listed CC oliseases associated with inappropriate expression of the proteins listed CC multifactorial diseases with a genetic component, such as autoimmune CC diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus exythromatosus and Grave's disease), inflammation, cancer CC leukaemia), diseases of the bladder, brain, breast, colon and kidney.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 22
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                                                               Matches
                                                                                               Query Match
                                                                                                                               Sequence 50 BP; 10 A; 14 C; 14 G; 12 T; 0 other;
                                                                                                                                                                       organisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 3552; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-465210/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shimkets RA, Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-DEC-1999; 99US-0173419
27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-DEC-2000; 2000WO-US35498.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W0200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL34304;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human SNP oligonucleotide #7512.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JAN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL34304 standard; DNA; 50 BP
                                                                             rocar
15 CCATTCTGCCTGGGTGACAATGCCTTGGTTTCAACACTATCGGAA 59
                                                               27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                53 ATCGGAATGTGACTGTCGGGCCTCCAGAAAACATTGA 89
                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9 ATCAAAATATGACAGTTTTATATCCATAAAATATTTA 45
                                                             Conservative
                                                                           17.48;
                                                           0;
                                                                                         Score 16.2; DB 22; Length 50;
                                                                           Pred.
                                                         Mismatches
                                                                             No.
                                                                           1.9e+04;
                                                         18;
                                                       Indels
                                                       0;
                                                     Gaps
                                                       0,
 YXX OS XXX
                                                                                                                                                                                                                                                                                                                                                                    В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 23
AAI64807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                             Cis-acting sequence; intron removal; trans-acting factor; alpha-subunit; RNA-activated protein kinase; eukaryotic initiation factor 2; eIF2alpha; tumour necrosis factor alpha; TNF-alpha; gene therapy; ss.
                 WO200014255-A1
                                                   Homo sapiens.
                                                                                                                                                           Cis-acting nucleotide sequence derived from human TNF-alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides the protein and coding sequences of staphylocoagulase repeated peptide 14. The sequences can be used in the treatment of cancer, haemopathy, HIV infection, immunological diseases and inflammation. The present sequence is a probe for the coding sequence
                                                                                                                                                                                               12-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                       AAZ99817 standard; RNA; 43 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 41 BP; 9 A; 10 C; 12 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide staphylocoagulase repeated peptide 14 and its encoding polynucleotide, useful for tumor, hemopathy, human immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-550467/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 7; Page 19(Disclosure); 31pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             virus infection, immunological disease and inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           мао Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CN1301711-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BODE-) BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Staphylocoagulase repeated peptide 14; cancer; haemopathy; HIV infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Staphylocoagulase repeated peptide 14 coding sequence probe #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAI64807;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI64807 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunological disease; inflammation; gene therapy; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            POCGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the invention.
                                                                                                                                                                                                                                                                                                                                                                                                  17 ATTCTGCCTGGGTGACAATGCCTT 40
                                                                                                                                                                                                                                                                                                                                                                  4 ATGCAGCCTGTGTGGGAATGCCTT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 CCATTGTGGCTAAGGTACAGTCCCGCAGGTTCACCAGGTTCTGCA 50
                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99CN-0125366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99CN-0125366.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        79.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17.28;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16; DB 22; Length 41; Pred. No. 2.1e+04;
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0 Gaps

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PN XXX XXX DEXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC capable of removing introns from a precursor transcript encoded by a ccapable of removing introns from a precursor transcript encoded by a ccapable of removal is effected during the production of mRNA of the gene, and cdepends on activation of a trans-acting factor which is an RNA-activated protein kinase capable of phosphorylating the alpha-subunit of eukaryotic circuit and the control of a cis-acting nucleotide sequence of the circuit control of a cis-acting nucleotide sequence of the circuit control of a cis-acting nucleotide sequence of the circuit control of a cis-acting nucleotide sequence of the circuit control of a cis-acting nucleotide control of the circuit control of a cis-acting nucleotide control of the circuit cis-acting nucleotide control of the circuit cis-acting nucleotide control of the circuit cis-acting nucleotide ci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-SEP-1998;
26-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kaempfer R, Osman F, Jarrous N, Ben-Asouli Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (YISS ) YISSUM RES & DEV CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 43 BP; 10 A; 13 C; 11 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The specification describes a cis-acting nucleotide sequence which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5; Page 15; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           alpha-subunit of eukaryotic initiation factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-257000/22.
                                                                                                                                                                                                                                                                     Cis-acting sequence; intron removal; trans-acting factor; alpha-subunit; RNA-activated protein kinase; eukaryotic initiation factor 2; eIF2alpha;
                                                                                                                                                                                                                                                                                                                                                                                           12-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ99815;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ99815 standard; RNA; 50 BP
  26-OCT-1998;
                                                                   06-SEP-1999;
                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                           Sequence of the stem loop of tumour necrosis factor-alpha gene.
                      07-SEP-1998;
                                                                                                                 16-MAR-2000
                                                                                                                                                                                                                                                      tumour necrosis factor alpha; TNF-alpha; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              68 TCGGGCCTCCAGAAAACATTGAGG 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 TGGGGCCTCCAGAACTCACTGGGG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98IL-0126112
98IL-0126757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-IL00483.
98IL-0126112
98IL-0126757
                                                                      99WO-IL00483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.1e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 21; Length 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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RESULT 26
AAL33971/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Regulation of gene expression by mRNA splicing is carried out using a cis-acting nucleotide sequence controlled by phosphorylation of the alpha-subunit of eukaryotic initiation factor 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; .2000-257000/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kaempfer R, Osman F, Jarrous N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 50 BP; 12 A; 15 C; 12 G; 11 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 7; Fig 5B; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (YISS ) YISSUM RES & DEV CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAL33971;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAL33971 standard; DNA; 50 BP
                                                                                  28-DEC-1999; 99US-0173419
27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                 05-JUL-2001.
                                                                                                                                                                                                                                                                                                                                                                   WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nervous system disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human SNP oligonucleotide #7179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JAN-2002 (first entry)
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                   28-DEC-2000; 2000WO-US35498.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11 UGGGGCCUCCAGAACUCACUGGGG 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     68 TCGGGCCTCCAGAAAACATTGAGG 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                     99US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17.2%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16; DB 21;
Pred. No. 2.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ben-Asouli Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 21; Length 50;
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Shimkets RA,

Leach M;

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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
         Claim I; Page 257; 475pp; English.
                                                       Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer,
                                                                                                                                                                           Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                      P-PSDB; AAM00634.
                                                                                                                                                                                                                                                 07-JAN-2000; 2000US-0174962.
                                                                                                                                                                                                                                                                                  05-JAN-2001; 2001WO-US00322
                                                                                                                                                                                                                                                                                                                                                               WO200151670-A2
                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; paternity test; forenein test; aberrant protein expression; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH89751 standard; DNA; 50 BP.
                                                 infection and diabetes -
                                                                                                                                                                                                                 (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                              19-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH89751;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 50 BP; 9 A; 21 C; 12 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoieti apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytokines, interferons, interleukins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 3446; 4143pp; English
                                                                                                                                        2001-451871/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  coding sequence polymorphic site SEQ ID NO: 532.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  49 GGCAGCACTCTGTGTAGGTGGCTGACGGTCCAGCCGA 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41 GGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCAGA 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                           Leach MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17.2%; Score 16; DB 22; Length 50; 62.5%; Pred. No. 2.2e+04; tive , 0; Mismatches 15; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8888888888%&
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                       acids in biological samples, and for giving progress in our understanding of Acquired Immunodeficiency Syndrome (AIDS). The primers are able to detect all HIV-1 and HIV-2 subtypes without detecting non-related viruses. The primer sets for HIV-1 and HIV-2 are compatible with each other, and can be combined to form a co-amplification assay for HIV-1 and HIV-2. Using more than one primer set to amplify target nucleic acid sequences which overlap a common probe region maximises
                                   strain sensitivity and robustness.
                                                                                                                                                                            The present PCR primer is used to amplify human deficiency type 2 (HIV-2) nucleic acids. The specification also describes primers and probes for HIV-1 and HIV-2. The primers and probes are useful for amplifying and detecting HIV-1 and HIV-2 and all their subtype nucleic
                                                                                                                                                                                                                                                                                 Disclosure; Page 11; 25pp; English.
                                                                                                                                                                                                                                                                                                                                     Detecting Human Immunodeficiency Virus 1 and 2 - using at least
                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-047891/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention.
                                                                                                                                                                                                                                                                                                                    four new oligonucleotide primers and multiple detection probes
                                                                                                                                                                                                                                                                                                                                                                                                                          Atwood SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ORTH-) ORTHO-CLINICAL DIAGNOSTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EP887427-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1; HIV-2; detection; Acquired Immunodeficiency Syndrome; AIDS; co-amplification assay; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV63698;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer used to amplify HIV-2 sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus type 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV63698 standard; DNA; 35 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 50 BP; 13 A; 18 C; 14 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           43 TGCCTGGGCGGCTGTGCCACAGTTTCTCCAGT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 TGCCTGGGTGACAATGCCTTGGTTTCAACACT 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                  Backus JW, Casey AE, Cummins TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97us-0050759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98EP-0304959.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.2e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                    Rasmussen EB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Sequence 35 BP; 6 A; 10 C; 5 G; 10 T; 4 other;

δõ

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RESULT 29
AAZ28217/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 뭉
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gene trapping; splicing; integration; mutation; genome; intron; exon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ28217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ28217 standard; DNA; 39 BP.
                                                                                                                                                                                                                                                       reading frame and/or internal ribosome entry site, a splice donor sequence and optionally intronic sequences. After integration of the sequence and optionally intronic sequences. After integration of the vector, transcription and splicing, chimeric RNAs are produced comprising the gene trap cassette spliced to an endogenous exon encoded within the target cell genome. Such gene trapping can be used to acquire novel sequence information from the trapped exons, useful to identify new genes and in screening e.g., to identify the genetic basis of diseases (such as cancer) or phenotypes, especially in the analysis of single nucleotide polymorphisms. The vector can also be introduced into a cell to activate the sequence of the control of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a murine splice acceptor sequence for use in a 5' gene trap cassette. Gene trapping can be used to simultaneously identify and murate genes. Gene trap vectors are designed to integrate into introns or genes in a manner that allows the cellular splicing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   splice acceptor; identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse splice acceptor sequence #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-DEC-1999 (first entry)
useful in screening, for example, to identify mutations associated with tumorigenic phenotypes. They can also be used to create cDNA libraries of cells, useful for large scale genetic analysis of the genome and to identify novel and mutated genes. The vectors are more efficient (e.g., their use identified 13-fold more genes) than conventional 3' gene trap vectors that rely on gene trapping as detected by antibiotic selection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarly a 3' cassette may comprise a promoter region, an exon (typically characterised by a translation initiation codon and open
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          machinery to splice vector-encoded exons to cellular mRNAs. vector may contain a 5' and/or a 3' gene trap cassette. A 5' cassette may comprise a strong splice acceptor sequence, sel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 13; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mutating genes in eukaryotic cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New gene trapping vectors, useful for identifying, activating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zambrowicz B, Friedrich GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (LEXI-) LEXICON GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-APR-1998;
14-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9950426-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              marker sequences and a polyadenylation site, but not a promoter.
                                                                                                                                                                                            expression of a naturally occurring cellular gene, useful to study gene function. They can be used to produce mutated cells or animals, and are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30 GTCSSSSCCAAAGAGAGAATTGAGGTG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           67 GTCGGGCCTCCAGAAAACATTGAGGTG 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 17.0%;
Similarity 63.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-0057328
98US-0081727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US06474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0079729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.8; DB 20;
Pred. No. 2.4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sands AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence, selectable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A gene trap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 30
                                  CA AAA29282-92 are examples of suitable splice acceptor sequences for use construction of genetically engineered vectors comprising 5, gene contragration into the host cell genome, enhance the number of cellular comprises a splice acceptor (SA), an exon sequence (EX1) located 3, to comprises a splice acceptor (SA), an exon sequence (EX1) located 3, to comprises a splice acceptor (SA), an exon sequence (EX1) located 3, to comprises a splice acceptor (SA), an exon sequence (EX1) located 3, to comprises a splice acceptor (SA), an exon sequence (EX1) located 3, end comprises a splice acceptor (SA), an exon sequence (EX1) located 3, end compressing the exon and a polyadenylation sequence defining the 3, end compressing the vectors may be used for gene discovery, gene cloning, gene committation, gene regulation, shuttling nucleic acid sequences throughout the genome and gene activation and over expression. The vectors can be cused to trap genes with a high level of efficiency regardless of whether compressed to trap genes with a high level of efficiency regardless of whether compressed to trap genes with a high level of efficiency regardless of whether compressed in the cell type into which the vector can be screened using the vectors, it is possible to produce large numbers of mutations and cover rapidly identify the mutated or trapped genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Use of vectors incorporating 5' gene trap cassettes increase the probability of identifying the 5' ends of gene open reading frames, important because these are difficult to obtain by conventional methods, and often code for the signal sequence in secreted and transmembrane proteins, an important group for potential drug targets.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 39 BP; 7 A; 17 C; 7 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Splice acceptor; vector; gene trap; gene discovery; cloning; analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA29286;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA29286 standard; DNA; 39 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-1998;
25-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               shuttling; gene activation; over-expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Splice acceptor sequence 5 for 5' gene trap cassette.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-400053/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (LEXI-) LEXICON GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cloning and gene mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zambrowicz B, Friedrich GA, Lilleberg S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 13; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Recombinant vectors for use in, for example, gene discovery, gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 17.0%;
Local Similarity 74.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   33 GGCGGTCCTGCAGGGAACAGTGGGGTG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           67 GTCGGGCCTCCAGAAAACATTGAGGTG 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0109302.
99US-0276533.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-US27366.

 Mismatches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.5e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sands AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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Query Match

Sequence 39 BP; 7 A; 17 C; 7 G; 8 T; 0 other;

17.0%; Score 15.8; DB 21;

Length 39;

В

Matches

20;

Conservative

0,

2.5e+04; -has 7; Indels

0;

Gaps

0

74.18;

Pred. No. Mismatches

Best Local Similarity

AAQ65503/c RESULT 32

AAQ65503 standard; DNA; 44 BP

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The invention relates to a translational repression system comprising a CC translational repressor operably linked to a constitutive promoter for use in cloning or expressing a specific heterologous gene. The CC use in cloning trapressor is bacteriophage MS2 coat protein. The improved cc expression vector systems are capable of expressing exogenous genes, in Escherichia coli and other host cells. The improved cc cincluding toxic genes, in Escherichia coli and other host cells. The control of the S2-based system can be used to stably clone and express cc certain toxic genes. The MS2-based 14 cassette is useful for the cloning of a category proteins that are useful in the production of a target control. The vector systems overcome the problem of promoter laskiness cc which leads to inappropriate transcription and expression of a gene cloning toxic genes. The transcriptional control proteins, e.g. moth and cc asiA of bacteriophage 14, regulate transcription and provide a staged conduction by acteriophage 14, regulate transcription and provide a staged conduction promoter system which is much less complicated and more transcription from specific promoters while inhibiting transcription cc transcription by inducing transcription for transcription by inducing transcription of protease genes. Sequences AAX89905-998 represent PCR primers used in the course of the cinvention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 31
AAX89906/c
ID AAX899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Š
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           Examples; Page 61; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A bacteriophage MS2 translational repression system for use in cloning or expressing specific heterologous genes, especially toxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-494095/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JAN-1999;
28-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Translational repression system; translational repressor; coconstitutive promoter; heterologous gene; bacteriophage MS2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9938985-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteriophage sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence of primer 909-35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer; protease gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      constitutive promoter; heterologous gene; bacteriophage MS2; t
transcriptional control protein; moth; asiA; bacteriophage T4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX89906;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX89906 standard; DNA; 40 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67 GTCGGGCCTCCAGAAAACATTGAGGTG 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33 GGCGGTCCTGCAGGGAACAGTGGGGTG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0072794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0237712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US01725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      74.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 2.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            coat protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          toxic gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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05-NOV-1992; (XYRO-) XYROFIN OY.

92US-0973325 93WO-FI00450.

05-NOV-1993;

11-MAY-1994. WO9410325-A. Synthetic.

Saccharomyces cerevisiae; ss.

Amplification; primer; polymerase chain reaction; PCR; carbon; xylitol; xylose; xylulose; transketolase;

Transketolase gene 3' primer (2271-2305).

30-NOV-1994 (first entry)

AAQ65503;

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17.0%; Score 15.8; DB 20; Length 40;
                                                                                                                                                                                                                                                                                                          밁
                                                                                                                                                                                                                                              RESULT 33
                                                                                                                                                                                                                                                                                                                                           Ş
                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
               Polynucleotide separation; ss; non-polar surface; STD multivalent cation free.
                                                                Polynucleotide separation method-related oligonucleotide (STD) SEQ ID #8.
                                                                                                                 27-SEP-2002 (first entry)
                                                                                                                                                              AAL43960;
                                                                                                                                                                                                 AAL43960 standard; DNA; 44 BP
                                                                                                                                                                                                                                                                                                                                                                                                         ocal
                                                                                                                                                                                                                                                                                                 39 GACCATTGCATTCTATAAGGGTGACAA 13
                                                                                                                                                                                                                                                                                                                                         8 GAGCACTCCATTCTGCCTGGGTGACAA 34
                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 74.18;
                                                                                                                                                                                                                                                                                                                                                                                   Conservative

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.8; DB 15; Pred. No. 2.5e+04;
                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     Length 44;
```

Sequence 44 BP; 11 A; 8 C; 10 G; 15 T; 0 other;

are given in AAQ65502-03

encoding transketolase was used to synthesise oligonucleotides for the cloning of this gene by PCR. The two oligonucleotides The known nucleotide sequence of Saccharomyces cerevisiae,

The two oligonucleotides

0

Gaps

0

D-xylulose or their oligomers

Example 9; Page 53; 90pp; English.

WPI; 1994-167479/20.

Apajalahti JHA, Harkki AM, Myasnikov AN, Pastinen OA;

Prodn. of xylitol from recombinant hosts - using a metabolic pathway which uses a carbon source other than D-xylose,

Query Match

Sequence 40 BP; 13 A; 9 C; 10 G; 8 T; 0 other;

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밁
                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           applying the mixture to a polymeric separation medium having non-polar surfaces. The non-polar surfaces are free from multivalent cations (preferably chromium, iron, nickel and/or copper) and are free to bind or interfere with the polynucleotides. The method of the invention is useful for separating a mixture that contains single-stranded polynucleotides (e.g. DNA and RNA) having lengths from 5 - 200 up to 2000 - 20000 nucleotides from impurities (e.g. failure sequences, salts and/or proteins). The method of the invention allows separation to be achieved within 10 - 30 minutes, in contrast to conventional gel chromatography which can require hours or days. The present DNA sequence represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Separation of a mixture of single-stranded polynucleotides e.g. DNA, or RNA, comprises applying the mixture to a polymeric separation medium having non-polar surfaces, and then separating the mixture \,\cdot\,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gjerde DT, Haefele RM, Hanna CP, Hornby D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-NOV-2000; 2000US-0714579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-SEP-2001; 2001WO-US27495.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 44 BP; 11 A; 11 C; 11 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention comprises a method of separating a mixture which contains single-stranded polynucleotides. The method of the invention involves
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TRAN-) TRANSGENOMIC INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAY-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200240130-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified.
                                                                                                                                                                    Cancer; treatment; vector; recombinase gene; transcription factor;
selective cell killing; gene therapy; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide that was used in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 24; Page 53; 91pp; English.
Hasegawa M,
                                                                       02-JUL-1998;
                                                                                                                                                                                                          WO 9909191 SeqID #5
                                                                                                                                                                                                                               28-MAY-1999 (first entry)
                       (DNAV-) DNAVEC RES INC
                                               20-AUG-1997;
                                                                                                25-FEB-1999
                                                                                                                     WO9909191-A1
                                                                                                                                                                                                                                                           AAX26503;
                                                                                                                                                                                                                                                                                  AAX26503 standard; DNA; 46 BP
                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                             35 CAAAGCATTGGTGGCATCTCCATCGGGACTCTACT 1
                                                                                                                                                                                                                                                                                                                                                                    32 CAATGCCTTGGTTTCAACACTATCGGAATGTGACT 66
                                                                                                                                                                                                                                                                                                                                                                                                     h 17.0%; score 15.8; DB 24
Similarity 65.7%; Pred. No. 2.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                            Conservative
Takeda K, Yokoi H;
                                                97JP-0223651.
                                                                       98WO-JP02993.
                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                   DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kuklin AI, Taylor PD;
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 44;
                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   밁
  RESULT 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ67047;
             map of the human genome
                       Novel biallelic markers used to construct a high density disequilibrium
                                                                       Cohen D,
                                                                                               (GEST ) GENSET.
                                                                                                                                   21-APR-1998;
                                                                                                                                                          21-APR-1999;
                                                                                                                                                                                  28-OCT-1999
                                                                                                                                                                                                           WO9954500-A2
                                                                                                                                                                                                                                                          variation
                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                        23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20;
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This invention describes a vector containing a recombinase gene which is controlled by (1) a promoter and action of which is in turn dependent on a specific transcription factor and (2) a desired gene to be expressed and two target sequences of the recombinase. Also described in the invention are (i) a host cell for introduction of the vector and (ii) and in vitro technique in which killing of cells without the specific transcription factor is selectively performed after the vector is introduced into a host cell by an in vitro technique. The technique is used for gene therapy e.g. in cancer treatment. The products of the invention allow the use of a specific vector specifically and safely, there is little toxicity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genomic map; haplotype; phenotype; polymorphic base; genotyping;
haplotyping; hybridisation; identification; characterisation;
diagnosis; single nucleotide polymorphism; SNP; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 46 BP; 12 A; 16 C; 8 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 12; 49pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gene expression specific to cells free from specific transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-181048/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human map-related biallelic marker SEQ ID NO:1394.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ67047 standard; DNA; 47 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                particularly in cancer treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         factor with constructed recombinase expression unit after infecting
WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genome; biallelic marker; high density disequilibrium map;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  40 TGTGAGCCAGGGCATTGGTTTAAACAC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25 TGGGTGACAATGCCTTGGTTTCAACAC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   - allowing expression of target gene in gene therapy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                              98US-0109732
                                                                                                                                                                                                                                            98US-0082614.
                                                                                                                                                                                                                                                                                                                     99WO-IB00822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                replace(24,G)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17.0%;
74.1%;
                                                                        Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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RESULT 36
AAA99938
ID AAA99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                               Ashkenazi AJ, Baker KP,
Goddard A, Gurney AL, Hi
Watanabe CK, Williams PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cardiovascular; endothelial; angiogenic disorder; pro179; pr0338; pr0354; pr0844; pr0846; pr01760; pr0205; pr0321; pr0333; pr0840; pr0877; pr0878; pr0879; pr0882; pr0885; pr0887;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ65554 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.
WPI; 2000-611444/58
                                                                                                                                                                                                                                                                                                                                                                  26-JUL-1999;
01-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                  02-JUN-1999;
20-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200053757-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide pl used for hybridisation analysis of DNA for PRO333.
                                                                                                                                                                                                         22-FEB-2000;
                                                                                                                                                                                                                                                                                          30-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy; oligonucleotide; ss.
                                                                                                                                                      (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-FEB-2000.;
                                                                                                                                                                                                                                                                                                                                         15-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JAN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA99938;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA99938 standard; DNA; 48 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 47 BP; 11 A; 12 C; 13 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ocal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11 GAAGCACTCAGGTATGCCTGAGTCACA 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7 GGAGCACTCCATTCTGCCTGGGTGACA 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                         2000WO-US04342
2000WO-US04414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000WO-US05004.
                                                                                                                                                                                                                                                              99WO-US28565.
                                                                                                                                                                                                                                                                                                                                                                                                                                        99WO-US12252.
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                                                                                                                                                                                                                                                                                                                                                                                         99US-0145698.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-US05028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                74.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17.0%;
                                                                           Hillan KJ,
                                                                     Ferrara N, Gerber H,
illan KJ, Marsters SA,
                                                     Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.8; DB 21; Pred. No. 2.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.6e+04;
                                                                     Gerritsen ME;
Paoni NF, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                     Pitti RM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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Novel PRO polypeptides and agonists and antagonists of them, used to diagnose and treat cardiovascular, endothelial and angiogenic disorders
                                                                                                    Example 19; Page 126; 181pp; English.
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Matches Query Match The present invention relates to methods for stimulating or inhibiting angiogenesis and cardiovascularization. The methods involve the use of pharmaceutical compositions based on the following proteins, PR0179, PR0238, PR0364, PR0844, PR0846, PR01760, PR0205, PR0321, PR0333, PR0840, PR0877, PR0879, PR0887, PR0885 or PR0887. These proteins were identified by isolating CDNA clones encoding secreted proteins. The proteins of the invention may be used to diagnose and treat cardiovascular, endothelial or angiogenic disorders. The present sequence is an oligonucleotide used for in situ hybridisation analysis Sequence 48 BP; 14 A; 16 C; 8 G; 10 T; 0 other; of one of the proteins of the invention. Local 16 TCACTATAGGGCCTCCACAGAACCTCG 42 62 TGACTGTCGGGCCTCCAGAAAACATTG 88 Similarity 74.18; Conservative 0; Mismatches Score 15.8; Pred. No. 2. .6e+04; DB 21; Indels Length 48; 0 Gaps

0

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RESULT 37
AAC58575
Human DNA41374 41374.pl probe SEQ ID NO:306.
                                        29-JAN-2001 (first entry)
                                                                                         AAC58575;
                                                                                                                                  AAC58575 standard; DNA; 48 BP
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Human; immune related disease; diagnosis; antiinflammatory; cardiant; dermatological; antiarthritic; antirheumatic; immunosuppressive; haemostatic; antiityroid; antiidabetic; nootropic; neuroprotective; antianaemic; hepatotropic; virucide; antisportatic; antialtergic; antianaemic; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; spondyloarthropathy; systemic sclerosis; sarcoidosis; diopathic inflammatory myopathy; Sjogren's syndrome; thyroiditis; systemic vasculitis; autoimmune haemolytic anaemia; diabetes mellitus; sutoimmune thrombocytopaemia; immune-mediated renal disease; demyelinating disease; hepatobiliary disease; Whipple's disease; inflammatory bowel disease; gluten-sensitive enteropathy; hybridisation; autoimmune disease; immune-mediated skin disease; allergic disease; numunological disease; transplantation associated disease; PCR primer; graft rejection; graft-versus-host-disease; probe; ss.

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Homo sapiens.
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ACC X ACC X

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WO200053758-A2
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14-SEP-2000

02-MAR-2000; 2000WO-US05841.

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10-MAR-1999;
12-MAR-1999;
23-MAR-1999;
12-APR-1999;
20-APR-1999;
28-APR-1999;
              14-MAY-1999;
02-JUN-1999;
                                       04-MAY-1999;
                                                                                                                              08-MAR-1999;
99US-0134287.
99WO-US12252.
99US-0141037.
                                     99WO-US08615.
99US-0131445.
99US-0132371.
                                                                                         99US-0125775
                                                                                                    99US-0123618
99US-0123957
                                                                                                                              99WO-US05028
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                            δ
                                                            Matches
                                                                                        Query Match
                                                                                                                                                                                          anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus, immune-mediated renal disease, demyelinating diseases of the central and peripheral nervous systems, hepatobiliary diseases, inflammatory bowel disease, gluten-sensitive enteropathy and Whipple's disease, autoimmune or immune-mediated skin diseases, allergic diseases, immunological diseases of the lung, and transplantation associated diseases including graft rejection and graft-versus-host-disease. AAC58397 to AAC58578 represent PCR primers and hybridisation probes used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-
30-NOV-
01-DEC-
01-DEC-
02-DEC-
02-DEC-
02-DEC-
02-DEC-
16-DEC-
20-DEC-
30-DEC-
30-DEC-
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                                                                                                                                                                                                                                                                                                                    proteins, anti-PRO antibodies, agonists and antagonists are useful for treating and diagnosing immune related disorders. The disorders are selected from systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis, idiopathic inflammatory myopathies, Sjogren's systemic vasculitis, sarcoidosis, autoimmune haemolytic syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sixty four PRO polypeptides, useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosis, rheumato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-SEP-
                                                                                                                      Sequence 48 BP; 14 A; 16 C; 8 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes sixty four human PRO proteins which can be used in the treatment of immune related diseases. The human PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 7; Page 151; 309pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         arthritis, osteoarthritis, thyroiditis and diabetes mellitus -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-572271/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stewart TA, Tumas D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kabakoff RC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-OCT-1999;
29-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-0CT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-SEP-1
                                                                                                                                                                 in the isolation of human PRO sequences. AAC58579 to AAC58642 and AAB33414 to AAB33477 represent human PRO polynucleotide and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ashkenazi AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JAN
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                                                                                                                                                    sequences given in the exemplification of the present invention.
                                                                           Local Similarity
16 TCACTATAGGGCCTCCACAGAACCTCG 42
                              62 TGACTGTCGGGCCTCCAGAAAACATTG 88
                                                            20;
                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000WO-US00219.
2000WO-US00277.
2000WO-US00376.
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2000WO-US04342.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baker KP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0144758.
99US-0145698.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US21547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-US21090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US20944
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                                                                       17.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pan J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Watanabe CK, Wood WI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Goddard A, G
                                                            0; Mismatches
                                                                         Score 15.8; DB 21
Pred. No. 2.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gurney AL, He
D, Shelton DL,
                                                                                     DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        erythematosis, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yan M;
                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hebert C, FDL, Smith V;
                                                                                      Length 48;
                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Henzel W;
                                                          Gaps
                                                            0;
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Considered numan OGHbetal4 protein may act as cell function of incated on human of the much of the protein may act as a cell function of intact human of the more of steroidogenesis or to inhibit the binding of intact human chromosome 14. This protein may act as cell function of intact human chromosome 14. This protein may act as a cell function of intact human chorionic gonadotropin (hog) to its receptor. The nucleotide sequence of the invention is useful to identify and obtain a concloride sequence of the invention is useful to identify and obtain a concloride sequence of the invention is useful to identify and obtain a concloride sequence of the invention is useful to identify and obtain a concloride progress of a patient of during drug therapy. An antibody against the human OGHbetal4 is useful conclored the properties of a patient during drug therapy. An antibody against the human OGHbetal4 is useful conclored human OGHbetal4 expression and monitor the progress of a patient during drug therapy.

Conclored human OGHbetal4 expression is implicated. The protein is conclored human OGHbetal4 expression is appecific human OGHbetal4 of including reproductive diseases useful for treating diseases or disorders including reproductive diseases useful for treating diseases or disorders including reproductive diseases useful for treating diseases or disorders conclored the invention such as thyroid deficiency or hyperactive thyroid contraception, blood and bone homeostasis. The present sequence represents the human OGHbetal4 cDNA specific real time TaqMan probe used to opuantitate expression of the human OGHbetal4 sequence of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK52192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated human OGHbetal4 polypeptide, useful in the treatment and diagnosis of reproductive diseases such as fertility, contraception, thyroid malfunction, altered cellular metabolism and bone homeostasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-435439/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OGHbeta14; human; ss; antiinfertility; contraceptive; antithyroid; chromosome 14; steroidogenesis; reproductive disease; fertility; contraception; pregnancy; menopause; dysmenorrhoea; thyroid malfunction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 31; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-OCT-2000; 2000WO-US27797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-OCT-2000; 2000WO-US27797.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200230978-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     thyroid deficiency; muscle atrophy; homeostasis; probe; TaqMan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human OGHbetal4 cDNA specific real time PCR probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK52192;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK52192 standard; DNA; 27 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (REGE-) REGENERON PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-APR-2002
                                            Invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "3' TAMRA labelled"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "6-FAM labelled"
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Sequence 27 BP; 9 A; 7 C; 7 G; 4 T; 0 other;

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                                                                                                                                           The present invention describes an N-type calcium channel deficient conn-human animal whose gene for the calcium channel has been disrupted. The gene that is disrupted encodes the N-type calcium channel alpha 1B subunit. Also described are: (1) a method for assaying usefulness of substances using the animal; (2) a method for screening for substances with potential pharmacological use; (3) useful substances found by the cut of the producting pharmaceuticals using this method (2) specifically methods for producing a hypotensive drug, a pain killer and a drug for lowering blood sugar and the substances themselves). The CC N-type calcium channel deficient non-human animal can be used for corrections substances for pharmaceutical use. Active substances include the present sequence represents a PCR primer used in the isolation of the mouse N-calcium channel alpha 1B subunit, which is used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 39
                                                       Matches
                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                           Sequence 30 BP; 4 A; 8 C; 11 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ino M, Miyamoto N, Niidome T, Sawada K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse; N-calcium channel alpha 1B subunit; blood pressure control; N-calcium channel knockout animal; blood glucose level control; pain transfer; hypotensive; analgesic; PCR primer; ss.
                                                                                                                                    example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                              N-type calcium channel deficient non-human animals useful for screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-OCT-1999; 99JP-0303809.
16-FEB-2000; 2000JP-0037839.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-AUG-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH21862 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 57; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (EISA ) EISAI CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-AUG-2000; 2000JP-0261979.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-OCT-2000; 2000WO-JP07503.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH21862;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200130137-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse N-calcium channel alpha 1B subunit PCR primer SEQ ID NO:5.
                                                                     госат
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FOCGE
                          71 GGCCTCCAGAAAACATTGAGGT 92
1 GGCCTCCAGGTCACAGTGAGGT 22
                                                      18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GAGCTGGGAGCACTCCATTCTG 22
                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           h 16.8%;
Similarity 81.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GACCTGGGAGAAACCCATTCTG 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                16.8%;
81.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Takahashi E, C
K, Nishizawa Y,
                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                  Score 15.6; DB 22; Length 30; Pred. No. 2.7e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.6; DB 24; Length 27; Pred. No. 2.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oki T, Yoshinaga T, Hatakeyama S;
I, Tanaka I;
                                                      4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4;
                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                    밁
                                                                                                                                                           Q
                                                                                                                                                                                                                                                                       ABL41371/c
                                                                                            RESULT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 40
AAI30206/c
                                                                                                                                                                                         Matches
                                                                                                                                                                                                                  Query Match
                                       ABL41371;
                                                                                                                                                                                                                                                                        testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI30206;
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of individuals were resequenced and single nucleotide polymorphisms (SNPs) in these genes discovered. The method is useful for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype. The nucleic acids containing the polymorphic sites may be useful in forensics and paternity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to the identification of nucleic acid molecules (AAI29513-AAI31314) from the human genome which include polymorphic sites which can predispose individuals to disease. Various genes from a number
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sites, useful in methods for predicting the presence, absence severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype \cdot
29-MAY-2002 (first entry)
                                                                                                                                 ABL41371 standard; DNA; 33 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 31 BP; 7 A; 6 C; 13 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 72; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecules from the human genome which include polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-522952/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cargill M, Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAY-2000; 2000US-0206129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAR-2001; 2001WO-US07268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200166800-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single nucleotide polymorphism; SNP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; resequence; genotype; disease; forensic; paternity testing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human single nucleotide polymorphism (SNP) SPTBN1 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI30206 standard; DNA; 31 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                              15 CCATTCTGCCTGGGTGACAATG 36
                                                                                                                                                                                                                                                                                                               22 CCGCTCTGCCTGGATGCCAATG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                    18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0187510.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              replace(16,T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lander ES;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.6; DB 22; Pred. No. 2.7e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                          AAZ07103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to human transcription aporepressor CCCTC-conjugated factor (CTCF) 10.23, the polyprucleotide encoding this polypeptide and DNA recombinant processes to produce the polypeptide. The present invention also discloses the application of the polyprucleotide for gene therapy. The present invention discloses the antagonist for resisting the polypeptide and its treatment effect. The present invention also discloses the method of applying the polypeptide in treating various diseases, such as common tumours, development disturbance disease. The present sequence is that of a PCR primer, useful in examples of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CN1326974-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human transcription aporepressor CTCF 10.23 PCR primer SEQ ID NO 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-JUN-2000; 2000CN-0116361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-2000; 2000CN-0116361.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polypeptide-human transcription arrestin CCCTC-conjugated factor (CTCF)10.23 and polynucleotide for encoding such polypeptide \dot{}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-217567/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 33 BP; 11 A; 8 C; 6 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 4; Page 19 (Disclosure); 34pp; Chinese.
             16-JAN-1998;
                                                                                                 27-JUL-1999
                                                                                                                                         JP11196888-A
                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                         aerobic coryneform bacterium; Brevibacterium flavum MJ-233;
                                                                                                                                                                                                                                                                                          PYC1; PYC1; PC gene; organic acid; pyruvic acid carboxylase;
                                                                                                                                                                                                                                                                                                                                      Recombinant PC gene PYC1 PCR primer b-4.
                                                                                                                                                                                                                                                                                                                                                                              07-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ07103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ07103 standard; DNA; 35 BP
                                                       16-JAN-1998;
                                                                                                                                                                              Saccharomyces cerevisiae
                                                                                                                                                                                                                                                PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Loca T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24 GAATGTGACTGTTGAGGATCCA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    57 GAATGTGACTGTCGGGCCTCCA 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transcription aporepressor; CCCTC-conjugated factor; CTCF;
herapy; tumour; development; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                          98JP-0020361.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.8%;
81.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.6; DB 24; Pred. No. 2.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ρy
AAV99100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             recombinant pyruvic acid carboxylase (PC) gene, where the PC gene can be derived from microorganisms, animals or plants, especially human, mouse, rat, yeast, or microorganisms of Corynebacterium, Bacillus, Rhizobium or Escherichia spp., or its extract with organic materials in a reaction solution containing carbonate ion or hydrogen-carbonate ion, or CO2, particularly prepared by addition of carbonic acid or hydrogen-carbonic acid or their salts, or CO2. The aerobic coryneform bacterium of Brevibacterium flavum, is preferably Brevibacterium flavum MJ-233 (PERM BP-1497). The method gives a high yield of organic acids (e.g. succinic acid) with an aerobic coryneform bacterium.

Brevibacterium flavum MJ-233, having a recombinant PC gene (e.g. PYC1 and PYC2). The present sequence represents a PCR primer for a recombinant PC gene (e.g. prombinant PC gene (e.g. prom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes the preparation of organic acids with the anaerobic action of an aerobic coryneform bacterium having a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 5; Page 10; 12pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        carbonate ion or hydrogen-carbonate ion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Preparing organic acids with a microorganism having a recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-471888/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 35 BP; 8 A; 9 C; 7 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and PYC2). The present sequence represent from of the present PC gene, designated PYC1, used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MITU ) MITSUBISHI CHEM CORP.
             New DNA methyltransferase nucleotide sequences - used particularly to develop antisense oligonucleotides for diagnostic and therapeutic purposes, particularly for inhibiting tumour growth
                                                                                                                                                                                                                                  17-DEC-1997;
30-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA methyltransferase genomic fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-MAR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV99100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV99100 standard; DNA; 43 BP.
                                                                                                                                                                                                                                                                                                                                                                                       W09854313-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         beta thalassemia; sickle cell anemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA methyltransferase; DNA MeTase; antisense oligonucleotide; human;
                                                                                                    WPI; 1999-059833/05
                                                                                                                                           Bigey P,
                                                                                                                                                                                     (UYMC-) UNIV MCGILL.
                                                                                                                                                                                                                                                                                                   29-MAY-1998;
                                                                                                                                                                                                                                                                                                                                             03-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14 ATGCCTTAGTTTCAACAGGAAC 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34 ATGCCTTGGTTTCAACACTATC 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                growth;
                                                                                                                                                Ramchandani S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                carboxylase gene - using reaction solution containing
                                                                                                                                                                                                                                    97US-0866340
                                                                                                                                                                                                                                                         97US-0069865
                                                                                                                                                                                                                                                                                                   98WO-IB01107.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour growth inhibition; silenced gene activation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16.8%;
                                                                                                                                                  Szyf M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.8e+04;
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Indels Length 35;

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Gaps

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AAZ68364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification
                                                                                                                      Novel biallelic markers used to construct a high density disequilibrium
                                                                         Claim 3; Page 808; 2745pp; English.
                                                                                                                                                                WPI; 2000-013267/01.
                                                                                                                                                                                                                                  (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                      21-APR-1999;
                                                                                                            map of the human genome
                                                                                                                                                                                                                                                                                    21-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                      WO9954500-A2.
                                                                                                                                                                                                                                                                    23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                       28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                         variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fragments. The sequences together sequentially produce the full length sequence ANY99128. The sequences can be used to derive antisense oligonucleotides which inhibit DNA MeTase expression. The antisense oligonucleotides can be used in investigating the role of DNA MeTase in cellular growth. They can be administered at different points in the cell cycle, or in conjugation with promoters or inhibitors of cell growth to determine the role of DNA MeTase in the growth of the cell type of interest. The antisense oligonucleotides can also be used for inhibiting tumour growth in a mammal, or to activate silenced genes to provide a missing gene function. This ameliorates disease symptoms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis; single nucleotide polymorphism; SNP; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genomic map: haplotype; phenotype; polymorphic base; genotyping;
haplotyping; hybridisation; identification; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human genome; biallelic marker; high density disequilibrium map;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human map-related biallelic marker SEQ ID NO:2711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 43 BP; 2 A; 6 C; 13 G; 22 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g. in beta thalassemia and sickle cell anemia. The antisense oligonucleotides can also be used an analytical and diagnostic tools
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ68364 standard; DNA; 47 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and a potentiators of transgenic plant and animal studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV99091-128 represent DNA methyltransferase (DNA MeTase) genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 44; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18 TTCTGCCTGGGTGACAATGCCTTGGTTTCA 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8
                                                                                                                                                                                                   Blumenfeld M,
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                                                                                                                                                                                                                                                                    98US-0082614.
98US-0109732.
                                                                                                                                                                                                                                                                                                                    99WO-IB00822.
                                                                                                                                                                                                                                                                                                                                                                                                                /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                     replace(24,T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16.8%;
70.0%;
                                                                                                                                                                                                 Chumakov I;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 43;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI73671;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treatment
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polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic
                                                                                                                                                                                                                                                                                                                                                        AA173060 to AA179867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 241; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-NOV-1999; 99US-0168138
29-NOV-2000; 2000US-0726173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-2000; 2000WO-US32758.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human silent SNP containing nucleic acid SEQ:612.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200140521-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               from the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    are not actually given a sequence in the Sequence Listing
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70.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 50 BP; 8 A; 16 C; 17 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                         g-protein beta3 subunit to predict physiological and pathological pathways in the human body. The invention may be used to detect, detect predisposition for, give a prognosis of, or treat a variety of diseases including high blood pressure, hypertonia, cardiac infarction, coronary disease and other heart or circulation diseases, diabetes, psychiatric diseases, immunological diseases, asthma, psoriasis, Crohn's disease, disease, immunological diseases, asthma, psoriasis, Crohn's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; G-protein subunit Gbeta3; variant; high blood pressure; asthma; hypertonia; cardiac infarction; coronary disease; heart disease; circulation disease; diabetes; psychiatric disease; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-FEB-2000; 2000DE-1004681.
21-FEB-2000; 2000DE-1007587.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-JUN-2000; 2000DE-1030945.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunological disease; psoriasis; colitis ulcerosa; transplant rejection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human G-protein subunit Gbeta3 intron 9 DNA fragment #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-OCT-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel use for sequence variants of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 3; 14pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of sequence variants of the beta3 subunit of human G protein to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-490007/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SIFF/) SIFFERT W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                  colitis ulcerosa, transplant rejection, HIV or Hepatitis B or C. This sequence represents a fragment of the human G-protein Gbeta3 subunit intron 9 which is used to illustrate the method of the invention.
                                                                                                               Sequence 50 BP; 14 A; 13 C; 19 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            predict,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ocal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunological and psychiatric diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detect, and determine drug targets for diseases including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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16.8%; Score 15.6; DB 22 70.0%; Pred. No. 3.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 50;
                                         Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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RESULT 47
AAH48693/c
                                                                          ABL50681/c
                                                                                                          RESULT 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel use for sequence variants of the human G-protein beta3 subunit to predict physiological and pathological pathways in the human body. The invention may be used to detect, detect predisposition for, give a prognosis of, or treat a variety of diseases including high blood pressure, hypertonia, cardiac infarction, coronary disease and other heart or circulation diseases, diabetes, psychiatric diseases, immunological diseases, asthma, psoriasis, Crohn's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of sequence variants of the beta3 subunit of human G protein to predict, detect, and determine drug targets for diseases including heart, immunological and psychiatric diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypertonia; cardiac infarction; coronary disease; heart disease; circulation disease; diabetes; psychiatric disease; Crohn's disease; immunological disease; psoriasis; colitis ulcerosa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; G-protein subunit Gbeta3; variant; high blood pressure; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ААН48693;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH48693 standard; DNA; 50 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JUN-2000; 2000DE-1030945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DE10030945-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human G-protein subunit Gbeta3 intron 9 DNA fragment #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-490007/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-FEB-2000; 2000DE-1007587.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transplant rejection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 3; 14pp; German.
ABL50681;
                                                     ABL50681 standard; DNA; 28 BP
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 50 BP; 15 A; 13 C; 18 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               intron 9 which is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       colitis ulcerosa, transplant rejection, HIV or Hepatitis B or C. This sequence represents a fragment of the human G-protein Gbeta3 subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SIFF/) SIFFERT W.
                                                                                                                                                                                                                                                                                                                             ocal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        40 CAGCCTCCCTGCCTGCCTCCCTTGGTTT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16 CATTCTGCCTGGGTGACAATGCCTTGGTTT 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21;
                                                                                                                                                                                          40 CAGCCTCCCTGCCTGTCCTTCCTTGGTTT 11
                                                                                                                                                                                                                                                16 CATTCTGCCTGGGTGACAATGCCTTGGTTT 45
                                                                                                                                                                                                                                                                                                                          h 16.8%;
Similarity 70.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000DE-1004681
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                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                        Score 15.6;
                                                                                                                                                                                                                                                                                                                             Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                              DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9
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                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                              Length 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                           0
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                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
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                                                                                                                                                                                                                                                                                                                                             RESULT 49
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                  01-CCT-1998;
                                                     09-MAR-2000
                                                                                  DE19845277-C1.
                                                                                                                                                            Wolframin; human; transmembrane protein; diagnosis; therapy;
                                                                                                                                                                                               Human wolframin intron 1/exon 2 boundary DNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G protein-coupled receptor (GPCR) proteins. The GPCR sequences have antidiarrheic and laxative activities. The GPCR sequences can be used for developing drugs to treat diseases of the digestive organs, e.g. colitis, diarrhoea, constipation and malabsorption syndrome, including gene diagnosis and therapy. The present sequence represents a PCR primer for rat GPCR, which is used in an example from the present
                                                                                                                     Homo sapiens
                                                                                                                                                     Wolfram Syndrome;
                                                                                                                                                                                                                                      26-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                             AAZ88883;
                                                                                                                                                                                                                                                                                                        AAZ88883 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 28 BP; 7 A; 11 C; 8 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes human and rat brain-originated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4; Page 71; 135pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human and rat brain-originated G protein-coupled receptor proteins and encoded DNAs, for developing drugs to treat diseases of the digestive organs, e.g. colitis, diarrhea, constipation and mal-absorption syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-269361/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-AUG-2000; 2000JP-0253862.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rat; rZAQ1; rZAQ2; G protein-coupled receptor; GPCR; antidiarrheic; laxative; drug development; digestive organ disease; colitis; diarrhoea; constipation; malabsorption syndrome; diagnosis; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-AUG-2001; 2001WO-JP07209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200216607-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rat G protein-coupled receptor protein PCR primer SEQ ID NO:30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oca1
                                                                                                                                                                                                                                                                                                                                                                                       26 CTATCAGCCTGTGGCTGCGGGGCCT 2
                                                                                                                                                                                                                                                                                                                                                                                                                          51 CTATCGGAATGTGACTGTCGGGCCT 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                            19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shintani Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                  98DE-1045277
                                                                                                                                             psychiatric disease; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16.6%;

    Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.4; DB 24; Pred. No. 3.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                            PR
PR
                                                                                                                                                                                                                                                                                                                                                                                XX OX XX DE
                                                                                                                                                                                                                                                                                                                                                                 PD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PA
XX
PI
XX
DR
XX
PI
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PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
of the matrix processing peptidase (MPP) was isolated by PCR
                A DNA sequence which codes for the mitochondrial targeting
                                                                               DNA encoding plant citrate synthase - used to regulate flower formation, to improve storage of tubers, etc. and to reduce sprouting
                                                     Example; Page 46; 87pp; English.
                                                                                                                                        WPI; 1995-328278/42.
                                                                                                                                                                           Mueller-roeber
                                                                                                                                                                                                                       (AGRE ) HOECHST-SCHERING AGREVO GMBH.
                                                                                                                                                                                                                                                                                                                              07-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mitochondrial targetting sequence; matrix processing peptidase;
                                                                                                                                                                                          La Cognata U,
                                                                                                                                                                                                                                                              22-SEP-1994;
                                                                                                                                                                                                                                                                             19-OCT-1994;
09-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                 14-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                 W09524487-A
                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-JAN-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT04205;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT04205 standard; DNA; 31 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               matrix processing peptidase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer oligo B for mitochondrial targeting sequence of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30 BP; 5 A; 8 C; 7 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wolfram Syndrome, in particular where there is a tendency to psychiatric disease. The products of the invention can be used for gene therapy. This sequence represents the intron 1/exon 2 boundary of human wolframin genomic DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel human transmembrane protein, wolframin wolframin and DNA encoding it are useful for diagnosis and/or therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 8; 26pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and therapy of Wolfram Syndrome, especially where there is a tendency to psychiatric disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wolframin, a transmembrane protein and related DNA useful for diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-184136/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Strom T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYMU-) UNIV MUENCHEN MAXIMILIANS LUDWIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30 GGAGTCCATCCTGCCTGGAAGAAAA 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10 GCACTCCATTCTGCCTGGGTGACAA 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Meitinger T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                      Landschutze V, Muller-rober B,
                                                                                                                                                                                                                                                          94DE-4435366
                                                                                                                                                                                                                                                                             94DE-4408629
                                                                                                                                                                                                                                                                                           94DE-4438821
                                                                                                                                                                                                                                                                                                                              95WO-EP00859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98DE-1045277.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16.6%;
76.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 3.2e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.4; DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6,
                                                                                                                                                                                      Landschuetze V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 30;
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RESULT 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pBluescript plasmid which contained the cDNA sequence of the MPP using the oligos AAT04204 and AAT04205. The resulting DNA fragment comprised the nts 299-397 of the sequence in Braun et al. 1992, EMBO J. 11:3219-3227, which codes for the MPP. An Asp 718 cleavage site was inserted at the 5' end of the sequence by oligo a. Oligo b inserted a BamHI site at the 3' end of the sequence.
                                                                                                                                                                                         to altered production of isoprenoids, steroids, flavonoids, hormones, fats, oils, rubber, alkaloids, sugars and starch and anti-pathogenic agents. A specific application is increasing the fat content (by at least 7%) of oilseed crops, e.g. (especially) rape, soya, sunflower and oil palms, but the method can also be applied to cereals, vegetables,
                                                                                                                                                                                                                                                                                                                                                  AAVI1379-V11390 are primers used in a method to construct a transgenic plant cell with altered acetyl-coenzyme A (AcCoA) hydrolase metabolism which has higher activity than wild-type cells. Controlling AcCoA hydrolase activity is used to regulate intracellular distribution of metabolites, specifically it increases concentration of acetate, leading
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Transgenic plants expressing heterologous acetyl-coenzyme A hydrolase - for increased enzymatic activity, has altered hydrolase of metabolites, specifically for oleaginous plants,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 31 BP; 7 A; 8 C; 9 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9806831-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV11384;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV11384 standard; DNA; 31 BP.
                                                                                                   Sequence 31 BP; 7 A; 8 C; 9 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 27; 65pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-159524/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Baeuerlein M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Solanum tuberosum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Acetyl Coenzyme A hydrolase; Acetyl-CoA; AcCoA; transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Potato matrix processing peptidase PCR primer mito-TP2
                                                                                                                                                                sugar beet, tobacco, cotton or ornamental plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         higher fat content in seeds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      isoprenoid; steroid; flavonoid; hormone; fat content; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                34 ATGCCTTGGTTTCAACACTATCGGA 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATCCCTTGGTTGCAACAGCAGCTGA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      La COGNATA U, Mueller-Roeber B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96DE-4032121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96DE-4033374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97WO-EP04311.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      intracellular ditribution;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.6%;
16.6%;

 Mismatches

Score 15.4; DB 19; Pred. No. 3.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.4;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      metabolite;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Willmitzer L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 31;
                            Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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                                                                                                                                                RESULT 53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                PCR primers AAV10419 & AAV10420 are used to amplify the potato matrix processing peptide (MPP) which is used in a in a method resulting in the production of transgenic plant cells with altered acetyl-CoA metabolism. The acetyl-CoA hydrolase activity of the cells is greater than that of wild-type cells as a result of the expression of a foreign DNA sequence encoding a protein with acetyl-CoA hydrolase activity. The transgenic plants may have one or more phenotypic traits selected from reduced or increased fatty acid levels in leaves, increased soluble sugar levels in leaves, increased starch levels in leaves, reduced growth, multiple shoot formation, and change in leaf colour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Acetyl-CoA-hydrolase; transgenic plant; phenotypic trait; altered metabolism; matrix processing peptidase; MPP; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Potato MPP PCR primer Mito-TP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV10420;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV10420 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-121465/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
Probe used to identify FtsY sequences in microorganisms
                                                                                                                                                                                                                                                                                                                                                    Sequence 31 BP; 7 A; 8 C; 9 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 10; 26pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Baeuerlein M, La Cognata U, Mueller-Roeber B, Willmitzer L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DE19632121-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Solanum tuberosum
                                    23-JUL-2001
                                                                        AAF89684;
                                                                                                              AAF89684 standard; DNA; 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transgenic plant cells – containing foreign acetyl-CoA hydrolase
                                                                                                                                                                                                                                                                                              Tocar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    34 ATGCCTTGGTTTCAACACTATCGGA 58
                                                                                                                                                                                                                                      34 ATGCCTTGGTTTCAACACTATCGGA 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 ATCCCTTGGTTGCAACAGCAGCTGA 31
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                                                                                                                                                                                                      7 ATCCCTTGGTTGCAACAGCAGCTGA 31
                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                            Conservative
                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96DE-1032121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96DE-1032121.
                                                                                                                                                                                                                                                                                              16.6%;
                                                                                                                                                                                                                                                                            0;

 Mismatches

                                                                                                                                                                                                                                                                            Score 15.4; DB 19;
Pred. No. 3.2e+04;
0; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6
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                                                                                                                                                                                                                                                                                                                Length 31;
                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  potato;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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RESULT 54
AAT65388
ID AAT653
XX
AC AAT65
XX
O9-SE
XX
O9-SE
XX
U9-SE
XX
U9-S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present probe was used to identify FtsY gene sequences from microorganisms. The probe is used in the method of the invention. The specification describes a method for identifying an organism among a population of organisms in a biological sample. The method involves amplifying a less conserved target sequence present between a pair of conserved regions, using primers corresponding to a pair of conserved regions in a genome of a population of organisms, and then using a probe to detect the amplicon. The method is used for identifying an organism e.g. a prokaryotic organism, bacteria, virus, or a single cell eukaryotic organism, among a population of organisms in a biological sample such as blood, urine, cerebrospinal fluid, sputum, tracheal aspirate or pleural fluid or in a tissue sample. The method is also useful for diagnosing a disease or disorder associated with an organism, and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       distinguishing the presence of bacteria and unicellular eukaryotes, or bacteria, yeast, paramecia, trypanosoma, unicellular eukaryotes or viruses. The method is useful for diagnosing bacterial sepsis and related septic shock associated with several bacterial infections, and for testing food-borne bacteria such as
                                      identification; SELEX; anti-mitogenic; inhibition; cell; Systematic Evolution of Ligands by Exponential enrichment; epithelial; proliferation; diagnosis; treatment; cancer;
                                                                                                                                                                                                                   Human keratinocyte growth factor binding ligand T15.
                                                                                                                                                                                                                                                                                                 09-SEP-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 39 BP; 6 A; 9 C; 10 G; 14 T; 0 other;
                                                                                                               Human; keratinocyte; growth factor; hKGF; binding ligand;
identification; SELEX; anti-mitogenic; inhibition; cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT65388 standard; RNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Escherichia coli and Salmonella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 33; Page 70; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying an organism from a population of organisms in a sample comprises amplifying a less conserved target sequence present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-NOV-1999; 99US-0165881.
06-JAN-2000; 2000US-0479457.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-355645/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-NOV-2000; 2000WO-US31579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200136683-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (APPO-) APPOLO BIOTECHNOLOGY INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Streptococcus pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bacterial infection; food-borne bacteria; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FtsY; microorganism identification; bacterial sepsis; septic shock;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Loca 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37 CCTTGGTTTCAACACTATCGGAATGTGACTGTC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 CCTTGGTGTCAATATTGACGAATTGCTCTTGTC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
   inflammatory bowel syndrome; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.6%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.4; DB 22; Pred. No. 3.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
   맑
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PR XXX PTT TXXX PR XXX PTT XXX PR XXX PTT XXX 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV51124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PF PR PR PR PR PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence, a human keratinocyte growth factor (hKGF) binding ligand, was identified by Systematic Evolution of Ligands by Exponential enrichment (SELEX). Briefly a candidate mixture of nucleic acids was contacted with hKGF, and nucleic acids having an increased affinity to hKGF partitioned from the remainder of the mixture. The partitioned nucleic acids were then amplified to yield a mixture of nucleic acids enriched for sequences with higher affinity and specificity for binding to hKGF. The ligand is anti-mitogenic and may be used to inhibit epithelial cell proliferation, or in the diagnosis and treatment of hKGF mediated and the contact of the sequences.
                                                         Polymorphic marker; allele-specific; primer; probe; amplification; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; maize; ss.
Zea mays.
                                                                                                                                                                                                                                                                                     11-JAN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                  AAV51124;
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV51124 standard; DNA; 41 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41 BP; 8 A; 10 C; 10 G; 13 U; 0 other;
                                                                                                                                                                                                                Maize polymorphic marker S27G3/G6-1B DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bowel syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 61; Page 173; 209pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gold L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pathological conditions, e.g. psoriasis, cancer and inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identification of nucleic acid ligands to TGF-beta, PDGF and hKGF using SELEX, used in the diagnosis and treatment of proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-034387/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-JUN-1995;
05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAR-1996;
02-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W09638579-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NEXS-) NEXSTAR PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30 GACAATGCCTTGGTTTCAACACTAT 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4 GACGAUGCGGUGGUCUCAAUUCUAU 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Janjic N, Pagratis N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96WO-US08014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95US-0479783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-0465594
95US-0479725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95US-0465591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95US-0458424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95US-0458423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-0618693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "all pyrimidines are 2'-fluoro modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16.6%;
52.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.4; DB 18; Pred. No. 3.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ringquist S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Toothman PJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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RESULT 56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This DNA sequence is a region of a Zea mays genome which contains a polymorphic marker. This sequence can be used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, on omnitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Landry BS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W09824796-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
  01-DEC-1997;
                                                                                                                                                                                                                                                                                                  AAV51123;
                                                                                                                                                                                                                                                                                                                           AAV51123 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 41 BP; 7 A; 7 C; 14 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 46; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brassica species allele-specific oligonucleotide probes and primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-JUN-1998.
                                                       W09824796-A1
                                                                                                                         variation
                                                                                                                                                                Zea mays
                                                                                                                                                                                           progeny; back-cross; hybrid; ancestry; maize; ss.
                                                                                                                                                                                                     Polymorphic marker; allele-specific; primer; probe; amplification; hybridisation; plant; hybrid certification; genetic contribution;
                                                                                                                                                                                                                                              Maize polymorphic marker S27G3/G6-1A DNA.
                                                                                                                                                                                                                                                                        11-JAN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AFFY-) AFFYMETRIX INC.
                             11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jocal
                                                                                                                                                                                                                                                                                                                                                                                                                      60 TGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                               8 TGTGACTGTGGGCCCGTGTGACAGCTTTTACGT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for plant breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lemieux B, Murigneux A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         96US-0032069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-0813507
  97WO-US21782.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97WO-US21782
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                                                                              /replace= "g"
/note= "polym
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /replace= "q"
                                                                                                                                  Location/Qualifiers
                                                                                                            '*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.6%;
                                                                                "polymorphism"

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.4;
Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sapolsky RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             .5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 57
AAF30319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В
  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
The present sequence is that of a primer used in the PCF
                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF30319;
                           Example 1; Page 50; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Flt-3 ligand; Fms-like tyrosine kinase; human; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This DNA sequence is a region of a Zea mays genome which contains a polymorphic marker. This sequence can be used in the construction of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brassica species allele-specific oligonucleotide probes and primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-MAR-1997;
02-DEC-1996;
Immunogenic compositions comprising Flt-3 ligand encoding polynucleotide and one or more antigen, or cytokine encoding polynucleotides, useful for suppressing tumour growth and for treating autoimmune diseases (e.g. rheumatoid arthritis)
                                                                                                                                                                                                                                          30-JUL-1999;
                                                                                                                                                                                                                                                                                 31-JUL-2000; 2000WO-US20679
                                                                                                                                                                                                                                                                                                                            08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunotherapy; therapy; tumour; cancer; lymphoma; gene therapy;
VR1605; cap independent translational enhancer; CITE; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cap independent translational enhancer PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF30319 standard; DNA; 44 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41 BP; 8 A; 7 C; 13 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 46; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-333252/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AFFY-) AFFYMETRIX INC.
                                                                                                           WPI; 2001-123319/13.
                                                                                                                                                       Hermanson GG;
                                                                                                                                                                                              (VICA-) VICAL INC
                                                                                                                                                                                                                                                                                                                                                                   WO200109303-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Landry BS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for plant breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60 TGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8 TGTGACTGTGGGCCCGTGTGACAGCTTTTACGT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lemieux B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0813507
96US-0032069
                                                                                                                                                                                                                                          99US-0146170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Murigneux A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sapolsky RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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AAZ68249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amplification of a cap independent translation enhancer (CITE) sequence from vector pCITE. The amplified product was used in the construction of generic cloning vector VR1605 (see AAF30339) containing constant regions of the human kappa light chain and gamma I heavy chain separated by the CITE sequence. This plasmid is used as a negative control for VR1623 (see AAF30315), a bicistronic chimeric idiotype vector. Coadministration of VR1623 and VR6200 (see AAF30313) encoding a secreted form (see AAB20187) of mouse Fms-like tyrosine kinase (Flt-3 ligand) demonstrated the
                 AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences, AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis
                                                                                                                                                                                                                                                                                                                                                                                                         21-APR-1998;
23-NOV-1998;
for disease states. Compositions and methods of the invention can also
                                                                                                                                                                              Claim 3; Page 784; 2745pp; English.
                                                                                                                                                                                                                        map of the human genome
                                                                                                                                                                                                                                         Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                      WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                             Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ68249;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ability of the invention to enhance protection from tumour challenge in a murine B-cell lymphoma model.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis; single nucleotide polymorphism; SNP; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human genome; biallelic marker; high density disequilibrium map;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human map-related biallelic marker SEQ ID NO: 2596.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ68249 standard; DNA; 47 BP
                                                                                                                                                                                                                                                                                                                                                                    (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 44 BP; 10 A; 13 C; 8 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22 GCCTGGGTGACAATGCCTTGGTTTCAACACTAT 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11 GCCTTGCTGGCCACGCGTTATTTTCCACCATAT 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                         Blumenfeld M, Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                         98US-0082614
98US-0109732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-IB00822.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          replace(24,T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.6%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.4; DB 22; Pred. No. 3.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
pk266_4 (see AAZ19899) from composite deposit ATCC 98700. The cwas originally isolated from a human foetal kidney cDNA library. It codes for a novel secreted protein (see AAY31835). The inven provides new human secreted proteins (see AAY31828-38) and polynucleotides (see AAZ1983-901) isolated from foetal cell, ad blood, adult brain and foetal kidney cDNA libraries. They are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data ar given. The polynucleotides are also stated to be useful for
                                                                                                                                                                             This oligonucleotide probe can be used to isolate cDNA clone
                                                                                                                                                                                                                 Disclosure; Page 105; 107pp; English.
                                                                                                                                                                                                                                                 potentially used as, e.g. vaccines
                                                                                                                                                                                                                                                                    New polynucleotides derived from murine foetal cell cDNA libraries,
                                                                                                                                                                                                                                                                                                      WPI; 1999-562059/47.
                                                                                                                                                                                                                                                                                                                                                                                                                              17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                              20-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W09947555-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Secreted protein; pk266_4; human; therapy; diagnosis; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human foetal kidney secreted protein clone pk266_4 probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-DEC-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ19908;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ19908 standard; cDNA; 29 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 47 BP; 18 A; 9 C; 7 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               effects from pharmaceutical agents acting on a disease as well as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side characterisation.
                                                                                                                                                                                                                                                                                                                                                                                           (GEMY ) GENETICS INST INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            POCST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34 ATGCCTTGGTTTCAACACTATCGGAATGTGACT 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9 ATGCCTCGGTTTCTTCACATAACAAATGTAAAT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                       мссоу лм,
                                                                                                                                                                                                                                                                                                                                      Treacy M,
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                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0078803
99US-0078803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US05939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "biotinylated phorphoramidite residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                                                                                                    LaVallie ER,
Agostino MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.4; DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                       Collins-Racie LA,
                                                                                                                                                                                                                                                                                                                                        Steininger RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 47;
                                                                                                                            The invention
                                                                                                                                                                                                                                                                                                                                                         Evans C;
                    data are
                                                                                                                                                            The clone
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                                                                                     Q
                             AAS17330
                                      RESULT 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                           Matches
                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 29 BP; 8 A; 3 C; 10 G; 7 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy.
                                                                                                                                                                 The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of crythropoletin, granulocyte colony stimulating factor protein and interferon alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hammerhead ribozyme #1561.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF01570;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF01570 standard; RNA; 29 BP.
AAS17330;
                                                                                                                                                                                                                                                              Claim 32; Page 91; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                   12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                       11-APR-2000; 2000WO-US09721.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200061729-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ribozyme; erythropoietin; granulocyte colony stimulating factor;
                   AAS17330 standard; DNA; 33 BP.
                                                                                                                                                  Sequence 29 BP; 11 A; 5 C; 5 G; 7 U; 1 other;
                                                                                                                                                                                                                                                                               useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoletin -
                                                                                                                                                                                                                                                                                                    Enzymatic and antisense nucleic acid inhibition of repressor genes,
                                                                                                                                                                                                                                                                                                                         WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 16.3%;
Local Similarity 85.0%;
                                                                                                                    Match 16.3%;
Local Similarity 69.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       34 ATGCCTTGGTTTCAACACTA 53
                                                                                      59 ATGTGACTGTCGGGCCTCCAGAAAACATT 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25 ATCCCTTAGTTTCAACACCA 6
                                                                     29 ATGTGACTTTCGNCTCATCAGTAGATATT 1
                                                                                                                                                                                                                                                                                                                                            Zwick M, Pavco P, McSwiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                    99US-0129390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                   Score 15.2; DB 21; Length 29; Pred. No. 3.7e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.2;
                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .7e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 29;
                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                      AAV46356
                                                                                                                               RESULT 62
                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200183530-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
Synthetic
                                                                                         AAV46356;
                                                                                                                                                                                                                                                                                                                                                                                                                                               periodontitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-041482/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CSLC-) CSL LTD.
                                                                                                                                                                                                    17;
                                                                                                                                                                                                              Similarity
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CC proteins PG32 and PG33, and the polyneptides derived from outer membrane CC proteins PG32 and PG33, and the polynucleotide sequences encoding CC them. The sequences of the invention can be used in compositions CC for raising an immune response in a subject against P. gingivalis and CC treating, preventing or reducing periodontitis or other P. gingivalis CC infections (e.g. gingivitis). Since periodontal disease has linkage with CC cardiovascular disease, the polypeptides are also useful for reducing CC the incidence or severity of cardiovascular disease or as an adjunct in CC treating the cardiovascular disease. An antibody raised against the CC compared to full-length proteins. AAS17314-AAS17337 represent multiple CC coverlapping oilgonicleotide primers used for the generation of PG32C cy high expression bias codon replaced DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Soluble Porphyromonas gingivalis polypeptide; PG32C; immune response; periodontitis; gingivitis; periodontal disease; cardiovascular diseas tooth paste; mouthwash; antibacterial; dental; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Overlapping primer #17 for P. gingivalis PG32C high expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel soluble polypeptides comprising fragment of Porphyromonas gingivalis PG32 and PG33 proteins, useful for eliciting immune response in subject against P.gingivalis, and for treating or preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to the isolation of soluble Porphyromonas gingivalis (PG) polypeptides derived from outer membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Page 34; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-APR-2001; 2001WO-AU00482
antitumour agent; PCR primer; ss
                                  Serine threonine kinase; VRK1; VRK2; B1R kinase; cell growth control;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 33 BP; 7 A; 6 C; 9 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-APR-2000; 2000AU-0007182
                                                                                                                PCR primer for serine threonine kinase.
                                                                                                                                                                                                  11-NOV-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                          AAV46356 standard; DNA; 36 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          39 TTGGTTTCAACACTATCGGA 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 TTGGTTTCAACGCTATTGAA 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Czajkowski L, Ross BC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2;
Pred. No. 3.
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                        WPI; 2000-619232/59
                                                  Roizman B, Ogle WO;
                                                                                                      09-APR-1999;
                                                                                                                                07-APR-2000; 2000WO-US09215.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence is a PCR primer for DNA encoding the serine threonine kinase of the invention. The two serine threonine kinase variants (VRK1 and VRK2; shown in AAW64771 and AAW64772 respectively) are highly expressed in actively growing cells (such as foetal organs), and have significant homology with the BIR kinase of vaccinia virus. The kinase can be used as a substrate for the screening of potential inhibitors, which can then be used in the control of cell growth and as antitumour which can then be used in the control of cell growth and as antitumour.
                                                                            (ARCH-) ARCH DEV CORP
                                                                                                                                                                                      WO200061794-A1
                                                                                                                                                                                                             Herpesvirus
                                                                                                                                                                                                                                             detection; identification; Usl.5; phosphorylation site; virucide;
                                                                                                                                                                                                                                                                                        Herpesvirus expression vector construction PCR primer P10.
                                                                                                                                                                                                                                       PCR primer; ss.
                                                                                                                                                                                                                                                              Herpesvirus; post translational modification; infection; inhibition;
                                                                                                                                                                                                                                                                                                                       14-FEB-2001
                                                                                                                                                                                                                                                                                                                                                 AAC64550;
                                                                                                                                                                                                                                                                                                                                                                         AAC64550 standard; DNA; 36 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 36 BP; 6 A; 9 C; 14 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 7; Page 38; 63pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Serine-threonine kinase highly expressed in actively growing cells - useful for development of cell growth inhibitors and antitumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-388133/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for control of cell growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CHUG-) CHUGAI RES INST MOLECULAR MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9829552-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                            μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 AGCTGGGAGCACTCCATTCTGCCTGGGTGACAATGC 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense DNA delivered via a suitable vector can also be used
                                                                                                                                                                                                                                                                                                                                                                                                                                       AGCTGCGGCCGGGTCTGCGGCTTAGGTGAAAATGC 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                      99US-0128500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96JP-0357864.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97WO-JP04855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.2;
Pred. No. 4e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QΥ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA168432/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
This invention describes a novel method for quantitative determination of nucleic acid (I) in a biological sample. A competitive, quantitative polymerase chain reaction (PCR) is performed using (i) forward and reverse primers that bind to a fragment of (I) suitable for specific detection; (ii) a known amount of competitor DNA (Ia), generated in an earlier PCR, that has the same size and base composition as (I) and differs only minimally from (I); (iii) usual PCR auxiliaries and (Iv) the sample. The individual strands of both amplicons (from (I) and (Ia)) are analyzed, directly or after a standard cleavage reaction, for mass and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAI68432;
                                                                                                                                                                                                                                                                                                                                                                                DE10001881-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
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Example 4.1; Page 9; 14pp; German.
                                                                                               Quantitative determination of nucleic acid, by competitive polymerase chain reaction in presence of known amount of competitor very similar
                                                                                                                                                                                                      WPI; 2001-603299/69.
                                                                                                                                                                                                                                                                                                                                                                                                         19-JAN-2000; 2000DE-1001881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Quantitative determination; high throughput screening; competitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 36 BP; 5 A; 12 C; 9 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           substances able to inhibit herpesvirus infections. The assay uses a polypeptide comprising a US1.5 phosphorylation site, a candidate inhibitor and a kinase and compares the phosphorylation state of the polypeptide with that of a similar polypeptide contacted with the kinase in the absence of the candidate inhibitor. The inhibitor of US1.5 phosphorylation is used to inhibit herpesvirus infection in a cell before or after the cell is contacted with herpesvirus. In particular the inhibitor is used to prevent herpes simplex virus infections. The present sequence represents a PCR primer used in
                                                                     to the target -
                                                                                                                                                                                                                                                                         Eschrich K, Rupf S;
                                                                                                                                                                                                                                                                                                                                           (UYLE ) UNIV LEIPZIG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JAN-2000; 2000DE-1001881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Methylobacterium organophilum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amplification; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     M. organophilum methanol dehydrogenase competitor primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI68432 standard; DNA; 36 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the construction of herpesvirus expression vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Assay for detecting inhibitors of herpesvirus infections monitors the phosphorylation state of polypeptides with a Usl.5 phosphorylation site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 91; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a screening assay to determine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64 ACTGTCGGGCCTCCAGAAAA 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28 AATGTCGGCCATCCAGAAAA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              υ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
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RESULT 65
AAV50773
Вb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA and DNA. The method is simple, quick, automatable and suitable for high throughput screening. The close similarity between sample and competitor DNA may eliminate the need for experimental confirmation of the efficiency of the amplification. This sequence represents a primer used in the amplification a nucleic acid fragment capable of acting as a competitor to the Methylobacterium organophilum methanol dehydrogenase (MDH) gene used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amount, so that they are unequivocally identified from mass and, from the ratio of their amounts and the known amount of (Ia), the amount of (I) is calculated. The method is used for quantitative determination of RNA and NNA mass matched is disclosed for quantitative determination of the product of the control of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 36 BP; 9 A; 10 C; 11 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brassica sp. polymorphic marker 85/18A11/86 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-JAN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV50773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV50773 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brassica sp
                                                                                                                                                                          This DNA sequence is a region of a Brassica napus or Brassica oleracea genome which contains a polymorphic marker. This sequence can be used in the construction of allele-specific primers and probes for
                                                                                                                                                                                                                                                                                                                                                           Brassica species allele-specific oligonucleotide probes and primers - useful for plant breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W09824796-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        progeny; back-cross; hybrid; ancestry; ss.
                                                  amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a
                                                                                                                                                                                                                                                                                                     Claim 1; Page 38; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hybridisation; plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic marker; allele-specific; primer; probe; amplification;
                       back-crossed plant with an ancestral plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-333252/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Landry BS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21 TGGGAGCCCTTCATGCTGCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 TGGGAGCACTCCATTCTGCC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lemieux B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96US-0032069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97WO-US21782
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /replace=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16.3%;
85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hybrid certification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murigneux A, Sapolsky RJ;

 Mismatches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No. 4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genetic contribution;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ω
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ABL96120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        So
                                                                                                                     XSXEFF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                     Query Match
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 41 BP; 10 A; 11 C; 8 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plant; polymorphism; polymorphic; single nucleotide polymorphism; SNP;
cruciferae; agriculture; food; canola; condiment; oilseed; vegetable;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brassica polymorphic site containing sequence 123-85/18A11/16.
                                                                                                                                                                                                                                                                                          The invention relates to oligonucleotides containing polymorphic sequences isolated from two Brassica species, B. napus and B. Oleracea, and a method for using them to identify, distinguish, and determine the relatedness of individual strains or pools of nucleic acids from plants within the family Cruciferae. This is useful because many cruciferous plants are important agricultural items and include many foodstuffs, for example, condiments, oilseeds, and vegetables. These condiments are incompleted to the sequence of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US6358686-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL96120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL96120 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brassica sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-AUG-2002 (first entry)
                                                                                                                     Sequence 41 BP; 10 A; 11 C; 8 G; 12 T; 0 other;
                                                                                                                                                                           include crops such as canola (a type of Brassica napus), which is one of the largest crops in Canada. The sequences given in records ABL96003-ABL96146 represent oligonucleotides containing polymorphic sequences isolated from B. napus and B. oleracea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic oligonucleotides from Brassica napus and B. oleracea useful for identifying, distinguishing, and determining the relatedness of individual strains or pools of nucleic acids from Cruciferae plants,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-314753/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lemieux B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Examples; Column 7-8; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 16.3%;
Local Similarity 63.9%;
                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28 GTGACAATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 GTGAAAAGGCATTGTTCTCCAAAATCTCGCTCTGCG 41
   23;
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Landry BS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96US-032066P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0464453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                          16.3%;
63.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     מ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sapolsky RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches 13;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.2;
Pred. No. 4.
                                 Score 15.2;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.2e+04;
                                    4.2e+04
                                                                   DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
      Gaps
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DP 68

28 GTGACAATGCCTTGGTTTCAACACTATCGGAATGTG 63

6 GTGAAAAGGCATTGTTCTCCAAAATCTCGCTCTGCG

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XXX DXXXX
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                                                                                                                                                                                                                                                                                                                               RESULT 68
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                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                 Query Match
                             Packaging expression cassette pCMVFTV17S construting primer FTV9.
                                                                                                                                                                                                                                                                                                                                               The present sequence is FIV9 primer used in the construction of packaging expression cassette pCMVFIVDelta17S. To generate pCMVFIVDelta 17S, in vitro mutagenesis was carried out using the primers FIV9 and FIV1 in the first round of PCR. The PCR products along with primers FIV1 and FIV4 were subjected to the second round of PCR, the product of which was digested with Sac I and Tthl11 I and ligated into similarly digested pF34NDeltaenv to generate pF34NDelta17Senv. pF34NDelta17Senv was cleaved with Sac I and ligated into XhOI/NotI digested pCMVFBeta to generate pCMVFIVDelta17S also referred to as pCFIVDeltaORF2.
Murin'e leukemia virus; MLV; feline immunodeficiency virus; FIV; chimeric;
                                                                   27-AUG-2002 (first entry)
                                                                                                 ABL60729;
                                                                                                                             ABL60729 standard; DNA; 46 BP.
                                                                                                                                                                                                                                                                                                                          Sequence 46 BP; 6 A; 20 C; 7 G; 12 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5C; Page 98; 170pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA sequences for gene therapy in transformed hosts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New feline immunodeficiency virus vectors containing heterologous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dubensky TW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9936511-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FIV9 primer; pCMVFIVDelta17S; packaging expression cassette; FIV1; mutagenesis; Sac I; Tth111 I; pF34NDeltaenv; pF34NDelta17Senv; FIV4; Not I; XhOI; pCMVBeta; pCFIVDeltaORF2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer FIV9 to generate pCMVFIVDelta17S, packaging expression cassette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-1999;
16-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX90642;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX90642 standard; DNA; 46 BP.
                                                                                                                                                                                                                                                                                    TOCS!
                                                                                                                                                                                                          13
                                                                                                                                                                                                                       13 CTCCATTCTGCCTGGGTGACAATGCCTTGG 42
                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                     CCCCATCCTACCTTGTYGACTGTCCCTCGG 42
                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hardy SF, Hsu D, Johnston JC, Sauter SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0231235
98US-0071731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9908-0114955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0086825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US01194.
                                                                                                                                                                                                                                                                               16.3%;
66.7%;
                                                                                                                                                                                                                                                                  <u>ب</u>
                                                                                                                                                                                                                                                                               Score 15.2; DB 20;
Pred. No. 4.4e+04;
                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                9,
                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                          Length 46;
                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                             0;
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AAZ66988 RESULT 69

AAZ66988 standard; DNA; 47

10-SEP-2001 (first entry)

AAZ66988;

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CC vector backbone and an FIV vector construct (1), comprising an mark of a desired heterologous sequence to a vertebrate or insect e.g. cat, to cat a desired heterologous sequence to a vertebrate or insect e.g. cat, to construct of the parameters of gene delivery in construct, and also useful in veterinary application. (1) is also useful in converse e.g. for stimulating a specific immune converses, inhibiting the interaction of an agent with a host cell receptor, to express a toxic palliative (conditional toxic palliatives), converse e.g. for stimulating a specific immune converse e.g. palliative (conditional toxic palliatives), converse e.g. converse
                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 46 BP; 6 A; 20 C; 7 G; 12 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               constructing a packaging expression cassette pCMVFIV17s.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric murine leukemia virus-feline leukemia virus vector construct, useful in gene therapy for treating diabetes, cystic fibrosis, cancer, hypercholesterolemia, hyperlipidemia, anemia, hemophilia, hepatitis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a chimeric murine leukemia virus (MLV)-feline immunodeficiency virus (FIV) vector construct (I), comprising an MLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 5; Page 89; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-471730/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dubensky TW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-NOV-2000; 2000US-253419P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic; virucide; hepatotrophic; antiinflammatory; antillpemic; antidiabetic; immunosuppressive; nootropic; neuroprotective; cardia antibacterial; fungicide; haemostatic; antianemic; anorectic; anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-NOV-2001; 2001WO-US44617.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200242482-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Feline immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy; mutagenic; PCR; primer; ss
                                                                                                                                                                                                                                                       LOCAL
13 CCCCATCCTACCTTGTYGACTGTCCCTCGG 42
                                                                                              13 CTCCATTCTGCCTGGGTGACAATGCCTTGG 42
                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gasmi M,
                                                                                                                                                                                                                                       16.3%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sauter SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           haemostatic; antianemic; anorectic; anti-HIV;

    Mismatches

                                                                                                                                                                                                                                                                                       Score 15.2; DB 24;
                                                                                                                                                                                                                                                  Pred. No.
                                                                                                                                                                                                                                                  4.4e+04;
                                                                                                                                                                                             9;
                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                       Length 46;
                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiant;
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                                                                                                                                                     RESULT 70
                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genomic map; haplotype; phenotype; polymorphic base; genotyping;
haplotyping; hybridisation; identification; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human genome; biallelic marker; high density disequilibrium map;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human map-related biallelic marker SEQ ID NO:1335.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ65654 to AAZ69578 represent human biallelic markers from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 515; 2745pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          map of the human genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis; single nucleotide polymorphism; SNP; ds.
               Plasmid pi-AN7 DNA/RNA hybrid oligonucleotide 15
                                                                                                                                                                                                                                                                                                                                                    Sequence 47 BP; 10 A; 7 C; 8 G; 22 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                    N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing
                                                                                                                                                                                                                                                                                                                                                                                                                                                         effects from pharmaceutical agents acting on a disease as well as other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-NOV-1998;
                                               13-DEC-1999 (first entry)
                                                                                    AAZ23367;
                                                                                                                      AAZ23367 standard; DNA; 48
                                                                                                                                                                                                                                                                                                                                                                                      from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                          treatment
                                                                                                                                                                                                                                                                                             Match 16.3%; Score 15.2; DB 21; Local Similarity 63.9%; Pred. No. 4.4e+04;
                                                                                                                                                                                                                                             13 CTCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAA 48
                                                                                                                                                                                                          12 CTCCTTTGGGGGAGATTGTCAATGTATTTTTTCAA 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blumenfeld M,
                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-0109732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0082614.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-IB00822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  replace(24,G)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chumakov I;
                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                Length 47;
                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                 0;
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XX PR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                XXXX
                                                                                                                                                                                                                                                                                                                                   밁
AAQ99206/c
                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          remove the mask and to generate an end that can react with further NA; and (5) repeating the cycle at least once using any suitable NA. The method is used to synthesize genes, regulatory elements, ribozymes etc., particularly those containing targeted mutations to alter their properties. The method can synthesize NA of any sequence or length, simply, inexpensively, and quickly, in either the 3'-5' or 5'-3' directions. It eliminates the need to store or transport sequences of gene size and allows in vitro gene mutation (to impart improvements in selected properties such as stability, solubility etc.). No intermediate cloning is necessary. AAZ23353-Z23367 represent DNA/RNA hybrid oligonucleotides derived from the plasmid pi-AN7 which are used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 misc_RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA/RNA hybrid; nucleic acid synthesis; in vitro gene mutation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          which comprises: (1) preparing a first NA (1) with at least one end that permits annealing and/or linkage of another NA; (2) annealing/linking the second NA to the end of (1), with the other end of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DE19812103-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes the novel synthesis of a nucleic acid (NA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 11; 20pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Iteratively synthesizing nucleic acid from oligonucleotides which contain cleavable sites, useful for preparation of genes, ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-541620/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-SEP-1999.
                 Polymerase chain reaction; PCR; primer; amplify; human; hypoxanthine-guanine-phosphoribosyl transferase; HPRT; human; probe; major histocompatibility complex; class I MHC antigen; transgenesis; immunodeficiency; transplant; ss.
                                                                                                                                                                                                                                   AAQ99206 standard; cDNA; 50 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 48 BP; 13 A; 10 C; 8 G; 16 T; 1 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the second NA masked; (3) masking the end of NA where the second NA has annealed; (4) cleaving the second NA at a predetermined site to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BERN/) BERNAUER A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAR-1998;
                                                                                                               Human HPRT gene probe primer PC31.
                                                                                                                                                          20-FEB-1996
                                                                                                                                                                                              AAQ99206;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                   48 GAAATTTGAGCGTCAGACCCCGTAGAAAAGATCAAAG 13
                                                                                                                                                                                                                                                                                                                                                                        56 GGAATGTGACTGTCGGGCCTCCAGAAAACATTGAGG 91
                                                                                                                                                                                                                                                                                                                                                                                                                 23;
                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98DE-1012103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "uracil"
                                                                                                                                                                                                                                                                                                                                                                                                                                    16.3%;
63.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2; DB 20;
Pred. No. 4.4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 48;
                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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0;

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RESULT 72
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                    13-JUN-2002
                                                 WO200246372-A1
                                                                             Synthetic
                                                                                                               ss; topoisomerase recognition site; topoisomerase; pcDNAGWDT(sc); pENTR-DT(sc); pcDNA-DEST41; pENTR/D-TOPO; pENTR/SD/D-TOPO; pcDNA3.2/V5/GWD-TOPO; pcDNA6.2/V5/GWD-TOPO; recombinational cloning;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antigen genes, creating non-immunoreactive cells. Cells containing inactivated MHC genes can be used as donors for transplantation, since they lack markers for host (recipient) immune attack. Transgenic mammals containing the inactivated MHC gene trait can be used in the study of immunodeficiency and may be used as a source of tissues and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         probe for a fragment of the human hypoxanthine-guanine-phosphoribosyl transferase (HPRT) gene. The probe was used in a southern blot on human retinal pigmented epithelial cell (RPE) DNA. The RPE's used had been targetted for a deletion of part of the HPRT gene. The HPRT fragment that was being searched for had the neomyvin resistance gene inserted into exon 3. The HPRT fragment with the insert was contained in the vector HPRT.MClneo.ro. This vector was introduced into RPE cells, and the relevant fragment of the HPRT gene was replaced by homologous recombination. The probe amplified by these two primers showed that this was the case. This indicates that this method can be used on MHC
                                                                                                                                                                            Adapter oligonucleotide TOPO 3.
                                                                                                                                                                                                          07-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                 ABK88813 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 50 BP; 16 A; 10 C; 8 G; 16 T; 0 other;
                                                                                                       gene targeting; mutation; TOPO 3; adapter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cells for transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genetically engineered mammalian cells lacking functional MHC antigens - are used to diminish immune attack when used for transplantation and as universal donor cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The sequences represented by AAQ99205-6 are amplification primers for a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 4; Page 34; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-246194/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Capon DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CELL-) CELL GENESYS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W09517911-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jourdes Arbones De Rafael M,
                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                     48 TCAATTCTCAGTGGTAGAAAATGCTTTG 21
                                                                                                                                                                                                                                                                                                                                                                14 TCCATTCTGCCTGGGTGACAATGCCTTG 41
                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dubridge RB,
                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93US-0175469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94WO-US14888.
                                                                                                                                                                                                                                                                                                                                                                                                           16.3%;
71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Greenburg G, Koller BH, Kucherlapati R;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.2; DB 16
Pred. No. 4.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Smithies O, Williams SR;
                                                                                                                                                                                                                                                                                                                                                                                                                      DB 16;
                                                                                                                                                                                                                                                                                                                                                                                             8
                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Length 50;
                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                       В
                                                                                                                                                                                                                                                                                                                               RESULT 73
Topoisomerase; recombination; ds
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CC topoisomerase recognition sites and/or one or more topoisomerases.

CR Also included are a vector comprising the nucleic acid, a vector chosen CC from pcDNAGMDT(sc), pENTR-DT(sc), pCNNA-DEST41, pENTR-DTOPO, CC a host cell comprising the nucleic acid or pcDNAS-ZVTS/GMD-TOPO or pcDNAS-ZVTS/GMD-TOPO, CC a host cell comprising the nucleic acid or vectors and an in vitro method CC of cloning a nucleic acid molecule involving: (a) obtaining a first CC nucleic acid molecule to be cloned; (b) mixing the first nucleic acid comprising at least a first topoisomerase recognition site flanked by at least a second recombination site, and at least a second recombination site, where CC cother, and at least one topoisomerase; and (c) incubating the mixture counter conditions such that the first nucleic acid molecule between the first and second recombination sites, thereby producing a first product conditions such that the first nucleic acid molecule between the first and second recombination sites, thereby producing a first product conditions and second recombination sites. The method is useful in methods for recombinational cloning and facilitates construction of gene ccomplete acid molecules or vectors which may be used to knockout cor mutate a sequence or gene of interest, particularly genes or sequences culcied cacid most culled used to make an or conditions and facilitate used to knockout cor mutate a sequence or gene of interest, particularly genes or sequences conclete acid most cells such as animal, plant, etc. Thus the animal conditions and facilitate used to make an or conditions and second for trangeting or mutating a sequence cor gene of interest, particularly genes or sequences conclete acid of to make an or conditions and cacid culties and conditions and cacid culties and conditions and cacid conditions are constructed to when a conditions and cacid conditions and cacid conditions and c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-DEC-2000; 2000US-254510P.
11-DEC-2000; 2000US-0732914.
14-SEP-2001; 2001US-318902P.
28-SEP-2001; 2001US-326092P.
27-NOV-2001; 2001US-333124P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of gene. The present sequence is an oligonucleotide used to make an adapter containing a consensus sequence for a topoisomerase site which may be included in the nucleic acid/vectors of the invention.
Topoisomerase adapter oligonucleotide, TOPO 3.
                                                                         18-JUN-2002 (first entry)
                                                                                                                                                                        ABK12602;
                                                                                                                                                                                                                                                           ABK12602 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 6 A; 4 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated nucleic acid molecule (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Page 216; 324pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               more topoisomerases, useful in recombinational cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated nucleic acid molecule comprises one or more recombination sites and one or more topoisomerase recognition sites and/or one or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-519662/55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brasch MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (INVI-) INVITROGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-DEC-2001; 2001WO-US45773.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CAACACTATCGGAAT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CAACACTATCGGAAT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cheo D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (a)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  one or more recombination sites; and (b) one or more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hartley JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leong L, man.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Madden K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 24; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Temple GF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gleeson M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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RESULT 74
                                                                                                                                                                                                                                                                                                                                                                                          В
                                                                                                                                                                                                                                                                                                                                                                                                                          QΥ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          topoisomerase covalently bound at or near a first end, and a second topoisomerase covalently bound at or near a second end, with a second ds nucleic acid molecule comprising a first blunt end and a second end. The method is useful for generating or constructing recombinant nucleic acids, and for covalently linking 2 or more nucleic acid molecules. The present method is simple and more efficient than previous methods for covalently linking nucleic acid sequences. The method greatly reduces the amount of work involved in screening to identify clones containing inserts in the desired orientation by enabling directional cloning efficiencies that are routinely in excess of 90%, and streamlines associated with the process. The present sequence represents a topoisomerase adapter oligonucleotide used in the methods of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to methods for generating a directionally linked recombinant nucleic acid molecule by contacting a topoisomerase-charged first double stranded (ds) nucleic-acid molecule comprising a first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-280935/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200216594-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
14-OCT-1999
                                                                        Synthetic.
                                                                                                                                                                                Expression plasmid construction oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 6 A; 4 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Fig 14; 116pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      molecule having 2 blunt ends -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         contacting a first double stranded (ds) nucleic-acid molecule with bound topoisomerases at both ends, with a second ds nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Generating directionally linked recombinant nucleic acid molecules by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-AUG-2000; 2000US-226563P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-AUG-2001; 2001WO-US26294
                                  WO9951766-A1.
                                                                                                                                                                                                                    11-JAN-2000 (first entry)
                                                                                                                                                                                                                                                            AAZ25796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SLOK ) SLOAN KETTERING INST CANCER RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INVI-) INVITROGEN CORP
                                                                                                       transcription factor; oncogene; tumour suppressor; ss.
                                                                                                                         simultaneous manipulation; expression vector; kinase;
                                                                                                                                       Yeast; ORF; open reading frame; PCR primer; library; expressible gene;
                                                                                                                                                                                                                                                                                             AAZ25796 standard; DNA; 16 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                            46 CAACACTATCGGAAT 60
                                                                                                                                                                                                                                                                                                                                                                                        1 CAACACTATCGGAAT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shuman S, Madden KR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.1%; Score 15; DB 24; Length 15; 100.0%; Pred. No. 3.6e+04;

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Неутап J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bennett RP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                         phosphatase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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ABK96427
ID ABK9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A method has been developed for producing a library of expressible coding regions. The method comprises: (a) amplifying coding regions using at least one coding region specific primer; (b) inserting each coding region into an expression vector; and (c) verifying the size and orientation of the inserted coding region. The method can be used for the rapid, simultaneous expression of large numbers of gene sequences. It can be used for producing expressible coding regions encoding human proteins such as kinases, phosphatases, transcription factors, oncogenes or tumour suppressors. The present sequence represents an oligonucleotide used in the exemplification of the present invention.
              Novel isolated fluorescent protein for producing fluorescence resonance energy transfer for use in in vivo labelling studies, is capable of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 16 BP; 7 A; 4 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Production of library of expressible coding regions, used for obtaining e.g. kinases, phosphatases, transcription factors, oncogenes or tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-APR-1998;
                                                               WPI; 2002-500276/53
                                                                                              Nys G,
                                                                                                                                                                                                                                          30-MAY-2002
                                                                                                                                                                                                                                                                          WO200242323-A2
                                                                                                                                                                                                                                                                                                        Discosoma sp.
                                                                                                                                                                                                                                                                                                                                                                     Coral; fluorescent protein; Anthozoa species; fluorescence emission;
                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer oGR48 for cDNA encoding Discosoma fluorescent protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                        24-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK96427;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK96427 standard; DNA; 29 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 86; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             suppressors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-620212/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fernandez JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (INVI-) INVITROGEN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-APR-1999;
                                                                                                                              (DEVG-) DEVGEN NV.
                                                                                                                                                            22-NOV-2000; 2000GB-0028495.
22-NOV-2000; 2000US-252790P.
                                                                                                                                                                                                           22-NOV-2001; 2001WO-EP13604.
                                                                                                                                                                                                                                                                                                                                      fluorescent labelling; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                     fluorescence absorbance; fluorescence resonance energy transfer; FRET;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46 CAACACTATCGGAAT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 CAACACTATCGGAAT 15
                                                                                              Plaetinck G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Heyman JA, Hoeffler JP, Marks-Hull HL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98US-0054936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-US07270.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16.1%; Score 15; DB 20;
100.0%; Pred. No. 3.7e+0.
htive 0; Mismatches
                                                                                            Bogaert T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.7e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sindici ML;
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emitting fluorescence upon irradiation by incident light

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RESULT 76
AAL425
XX
AAL425
XX
AAL425
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O8-JUL
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O8-JUL
XX
HUMAN
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HUMAN
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HOMO S
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CN1329
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CN1329
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Š
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
The invention comprises the amino acid and coding sequence of the human dynein receptor similarity kinase 51.26. The dynein receptor similarity kinase, 51.26 DNA and protein sequences are useful for treating tumours,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a label and/or marker, in particular as a genetic marker and/or expression marker, in the fields of (micro-) biology, biochemistry and/or molecular biology. They are also useful for monitoring expression of proteins within biological systems and subcellular localisation or trafficking of proteins. The polynucleotide sequences encoding the fluorescent proteins are useful for in vitro applications such as hybridisation and/or immunological assays. They are also useful for labelling polypeptides of interest.

ABK96395-ABK96461 represent PCR primers used in the examples of the present inventor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to novel coral fluorescent proteins isolated from two brightly fluorescent Anthozoa species (Polythoa and Discosoma species), and the polynucleotide sequences encoding them. The fluorescent proteins of the invention are capable of emitting fluorescence upon incident light irradiation, where the maximal absorbance of the incident light irradiation mm, and maximal fluorescence emission is 470-510 mm. The fluorescent proteins are useful for producing fluorescence resonance energy transfer (FRET), between a denor and accept fluorescence resonance energy transfer (FRET).
                                                                                                                                                        Example 4; Page 20 (Disclosure); 36pp; Chinese.
                                                                                                                                                                                                                        New polypeptide-human dynein receptor similarity kinase 51.26 for treating tumors, embryonic developmental deformity, various inflammations and various orthopedic diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mao Y, Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUN-2000; 2000CN-0116624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; PCR; primer; dynein receptor similarity kinase 51.26; tumour; embryonic developmental deformity; inflammation; orthopaedic disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human dynein receptor similarity kinase 51-26 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAL42596;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAL42596 standard; DNA; 33 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 29 BP; 7 A; 8 C; 7 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SHAN-) SHANGHAI BIODOOR GENE DEV CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JUN-2000; 2000CN-0116624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 between a donor and acceptor fluorescent protein molecule. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 33; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fluorescent proteins are useful for in vivo labelling studies, as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Toca T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 66 TGTCGGGCCTCCAGAAAACATTG 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 TGTCAGGCATCCCGAAGACATCG 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.1%;
78.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15; DB 24;
Pred. No. 4.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 77
CC translational repressor operably linked to a constitutive promoter for CC translational repressor operably linked to a constitutive gene. The constitutive promoter for CC use in cloning or expressing a specific heterologous gene. The improved CC translational repressor is bacteriophage MS2 coat protein. The improved CC expression vector systems are capable of expressing exogenous genes, cc including toxic genes, in Escherichia coli and other host cells. The CC bacteriophage MS2-based system can be used to stably clone and express CC control toxic genes. The MS2-based T4 cassette is useful for the cloning cof accessory proteins that are useful in the production of a target CC protein. The vector systems overcome the problem of promoter leakiness CC which leads to inappropriate transcription and expression of a gene CC cloning toxic genes. The transcriptional control proteins, e.g. moth and CC cloning toxic genes. The transcriptional control proteins, e.g. moth and CC asiA of bacteriophage T4, regulate transcription and provide a staged inducible promoter system which is much less complicated and more versatile than prior art non-staged systems. T4 middle promoters direct transcription from specific promoters while inhibiting transcription for from E. coli promoters, which therefore minimises competition for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Translational repression system; translational repressor; coat protein; constitutive promoter; heterologous gene; bacteriophage MS2; toxic gene transcriptional control protein; moth; asiA; bacteriophage T4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   embryonic developmental deformity, inflammations and various orthopaedic diseases. The present DNA sequence represents a PCR primer that is specific for the gene sequence of the human dynein receptor similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Examples; Page 61; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cloning or expressing specific heterologous genes, especially toxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A bacteriophage MS2 translational repression system for use in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-494095/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brown WC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JAN-1999;
28-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX89909 standard; DNA; 36 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 33 BP; 9 A; 8 C; 9 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9938985-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence of primer 909-38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       kinase 51.26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteriophage sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; protease gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        32 AGCCCTCCATTCTTCGGGTATGAGGATCCAT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9 AGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-US01725.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-0072794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0237712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ŏ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 33;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       toxic gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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             Creaming the invention relates to a translational repression system comprising a carranslational repressor operably linked to a constitutive promoter for cuse in cloning or expressing a specific heterologous gene. The cranslational repressor is bacteriophage MS2 coat protein. The improved cuse in cloning or expressing a specific heterologous gene. The curvession vector systems are capable of expressing exogenous genes, in Escherichia coli and other host cells. The charteriophage MS2-based system can be used to stably clone and express correction toxic genes. The MS2-based T4 cassette is useful for the cloning coff a casessory proteins that are useful in the production of a target correction. The vector systems overcome the problem of promoter leakiness cull neads to inappropriate transcription and expression of a gene coloning toxic genes. The transcriptional control proteins, e.g. mota and conductible promoter system which is much less complicated and more versatile than prior art non-staged systems. T4 middle promoters direct cranscription from specific promoters while inhibiting transcription for complete control promoters, which therefore minimises competition for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          translational apparatus and inhibits the cell from responding to target protein production by inducing transcription of protease genes. Sequences AAX89905-998 represent PCR primers used in the course of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Translational repression system; translational repressor; coat protein; constitutive promoter; heterologous gene; bacteriophage MS2; toxic gene transcriptional control protein; motA; asiA; bacteriophage T4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 36 BP; 8 A; 8 C; 8 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                       Examples; Page 61; 133pp; English.
                                                                                                                                                                                                                                                                                                                                                                    A bacteriophage MS2 translational repression system for use in
                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-494095/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Brown WC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9938985-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer; protease gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence of primer 909-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX89910;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX89910 standard; DNA; 36 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bacteriophage sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        35 GAGTTGGAAGAAATCCATTCAGC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GAGCTGGGAGCACTCCATTCTGC 23
                                                                                                                                                                                                                                                                                                                                                  expressing specific heterologous genes, especially toxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0072794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0237712.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-US01725.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.1%;
78.3%;

 Mismatches

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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.8e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               toxic gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             $8888
                                                                                                                                   RESULT 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ99700/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                         Matches
                                                      Query Match
                                      Best Local Similarity
                                                                                      Sequence 41 BP; 13 A; 12 C; 3 G; 13 T; 0 other;
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isolation of the bovine respiratory syncytial virus glycoprotein gene. This gene is used in the construction of homology vector 727-20.5. New recombinant swinepox (SPV) viruses have been made which contain a foreign DNA sequence inserted into a non-essential site of the SPV genome via the use of an homology vector. Homology vector 515-85.1 is a plasmid useful for the insertion of foreign DNA into the SPV genome. The plasmid 515-85.1 contains a unique Accl restriction site into which foreign DNA may be cloned. Recombinant SPVs can be used for immunising an animal against e.g. a human pathogen or equine or bovine pathogen, such as equine influenza A, equine herpes virus, bovine respiratory syncitial virus, bovine parainfluenza virus, infectious bursal disease virus or bovine diarrhoea virus. The recombinant viruses are also useful for enhancing a human immune response. The viral products can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ99700;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       translational apparatus and inhibits the cell from responding to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 77; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New recombinant swinepox viruses - having foreign DNA inserted into a site which is non-essential for viral replication, used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1995-075025/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cochran MD, Junker DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9503070-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bovine respiratory syncytial virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Swinepox; vaccine production; genetic engineering; recombinant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bovine respiratory syncitial virus glycoprotein gene PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-FEB-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ99700 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 36 BP; 11 A; 9 C; 10 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein production by inducing transcription of protease genes.
also be used for antibody production and in disease diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vaccine prodn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SYTR ) SYNTRO CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JUL-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vector; antigen; transcription; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAX89905-998 represent PCR primers used in the course of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GAGCTGGGAGCACTCCATTCTGC 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 GAGTTGGAAGAAATCCATTCAGC 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         930S-0097554.
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78.3%;
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Pred. No. 4.8e+04;
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Conservative

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16.1%; 67.7%;

Score 15; DB 16 Pred. No. 5e+04; Mismatches

DB 16; Length 41; 10; Indels

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Gaps

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RESULT 81
AAV47871
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       14-003-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The attachment (G) gene coding region from the bovine respiratory syncytial virus strain 375 (ATCC VR 1339) was cloned using a primer (AAQ97659) for cDNA priming and combined with a primer (AAQ97660) for PCR. The nucleocapsid (N) gene coding region was similarly cloned using primers AAQ97657 and AAQ97658, and the fusion (F) gene using primers AAQ97657 and AAQ976658, and the fusion (F) gene using primers AAQ97661 and AAQ97662. The PCR products were used to construct novel, foetal-safe, live recombinant infectious bovine rhinotracheitis virus vectors, e.g. S-IBR-052 (ATCC VR 2443), attenuated by deletion of the US2, gG and/or gE gene regions and also including foreign DAN coding for other infectious disease pathogen antigens. The recombinant vaccine
                                                                          AAV47871;
                                                                                                                                       AAV47871 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41 BP; 13 A; 12 C; 3 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel recombinant infectious bovine rhinotracheitis virus - used vaccines to immunise animals against bovine respiratory disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-FEB-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-APR-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ97660 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 84; 301pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-283781/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cocnran MD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bovine respiratory syncytial virus attachment gene PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ97660;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (shipping fever).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SYTR ) SYNTRO CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Loca1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used to protect cattle against bovine respiratory disease complex
                                                                                                                                                                                                                                                                                                       40 AATGTCTTGAATTTAAGATGATGGGTATGAG 10
                                                                                                                                                                                                                                                                                                                                                                   33 AATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            40 AATGTCTTGAATTTAAGATGATGGGTATGAG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     33 AATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94US-0191866
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.1%; Score 15; DB 16 67.7%; Pred. No. 5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 16; Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                   10;
                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     distinguish between strains. The polymorphisms may correlate with phenotypic traits (e.g. plant growth rate or crop yield), and the segments are useful to determine the presence/absence of specific polymorphisms correlating with the existence/absence of particular traits. The segments are also useful in marker assisted back-cross techniques to select plants with a higher percentage of recurrent parent in a back-cross population. Segments incorporate SNPs which occur more frequently than other polymorphism types and are therefore more likely to be located close to genetic loci of interest; different forms of characterised SNPs are also often easier to detect than other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a nucleic acid segment comprising at least 10 contiguous nucleotides from a vegetal sequence including a polymorphic site which is a single nucleotide polymorphism (SNP), or the complement of the segment. Also described are: (1) an allele-specific oligonucleotides hybridising to segment, or their complements, and (2) a method of analysing nucleic acids from a subject, by determining if a base is occupying any one (or a set) of polymorphic sites in 261 as segments are useful in fingerprint analysis in plants to determine which polymorphisms are present, which strain a plant belongs to and to polymorphisms are present, which strain a plant belongs to and to
Maize; marker; probe; PCR primer; polymorphism; vegetal sequence;
                                                                                                                                                                                                   AAV47872;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 41 BP; 8 A; 6 C; 13 G; 13 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vegetal sequences including single nucleotide polymorphism - useful, e.g. to determine polymorphisms in plants, determine strain in plant
                                                            Maize polymorphic site oligonucleotide marker UMC65-G3/G6-1B.
                                                                                                                            14-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                AAV47872 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 14; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-399160/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphism types.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     breeding and to correlate polymorphisms with phenotypic traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murigneux A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BIOC-) BIOCEM SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9830717-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lea sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphic site; corn; graminae species; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Maize; marker; probe; PCR primer; polymorphism; vegetal sequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maize polymorphic site oligonucleotide marker UMC65-G3/G6-1A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.18;
Local Similarity 63.68;
nes 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60 TGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                8 TGTGACTGTGGGCSCGTGTGACAGCTTTTACGT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96US-0032069.
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    Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15; DB 19
Pred. No. 5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 41;
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Gaps

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OS XX
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                                                                                                                                                                                                                                         RESULT 83
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonuclectides hybridising to segment, or their complements, and (2) a method of analysing nucleic acids from a subject, by determining if a base is occupying any one (or a set) of polymorphic sites in 261 sequences derived from is maize lines (see ANV47701 to AAV47961). The segments are useful in fingerprint analysis in plants to determine which polymorphisms are present, which strain a plant belongs to and to distinguish between strains. The polymorphisms may correlate with phenotypic traits (e.g. plant growth rate or crop yield), and the segments are useful to determine the presence/absence of specific segments are useful to determine the presence/absence of specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphisms correlating with the existence/absence of particular traits. The segments are also useful in marker assisted back-cross techniques to select plants with a higher percentage of recurrent parent in a back-cross population. Segments incorporate SNPs which occur more frequently than other polymorphism types and are therefore more likely to be located close to genetic loci of interest; different forms of characterised SNPs are also often easier to detect than other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a nucleic acid segment comprising at least 10 contiguous nucleotides from a vegetal sequence including a polymorphic site which is a single nucleotide polymorphism (SNP), or the complement of the segment. Also described are: (1) an allele-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vegetal sequences including single nucleotide polymorphism - useful, e.g. to determine polymorphisms in plants, determine strain in plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-399160/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BIOC-) BIOCEM SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-JUL-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9830717-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polymorphic site; corn; graminae species;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 2; Page 14; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               breeding and to correlate polymorphisms with phenotypic traits
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 41 BP; 7 A; 6 C; 14 G; 13 T; 1 other;
Synthetic
                                 pseudorabies virus; feline immunodeficiency virus; FIV; heartworm;
Dirofilaria immitis; PCR primer; ss.
                                                                   Swinepox virus; SPV; recombinant; vaccine; immunisation; diagnosis;
                                                                                                     BRSV strain 375 (VR-1339) G gene PCR primer from WO9804684.
                                                                                                                                                                         AAV26221;
                                                                                                                                                                                                         AAV26221 standard; DNA; 41 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphism types.
                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                               60 TGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                             8 TGTGACTGTGGGCSCGTGTGACAGCTTTTACGT 40
                                                                                                                                                                                                                                                                                                                                                               21; Conservative
                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9605-0032069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97WO-EP07134.
                                                                                                                                                                                                                                                                                                                                                                                  16.1%;
                                                                                                                                                                                                                                                                                                                                                                                  Score 15; DB 19
Pred. No. 5e+04;
                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                   DB 19; Length 41;
                                                                                                                                                                                                                                                                                                                                                                     11;
                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Вb
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Query Match
                                                                                                                                                                                                                                                The present sequence represents a PCR primer from the present invention. The present invention specifically describes recombinant swinepox virus (SPV) comprising a foreign DNA (I) inserted into a SPV genome which is capable of being expressed in a host cell into which the virus is introduced, where (I) is inserted into: (a) an ECORI site within a region corresponding to a 3.2 kb subfragment of the HindIII K fragment which contains both a HindIII and an ECORI site, of the SPV genome, and optionally (b) an ACCI site within a region corresponding to a 3.6 kb HindIII to BgilI subfragment of the HindIII M corresponding to a 3.6 kb HindIII to BgilI subfragment of the HindIII M fragment. The recombinant SPV can be used in a vaccine for immunising an animal against SPV. The invention also provides a method for testing a swine to determine whether the swine has been vaccinated with the
                                                                                                                                    vaccine, particularly containing S-SPV-008, or is infected with a naturally occurring wild-type pseudorables virus. Also (I) inserted into recombinant SPV can be used in a diagnostic assay, e.g. Feline immunodeficiency virus (FIV) env and gag genes and Dirofilaria immitts pag and 22kd are useful to detect feline immunodeficiency caused by FIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9804684-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 49; 473pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    animal against swine pox virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SYTR ) SYNTRO CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JUL-1996;
                                                                     Sequence 41 BP; 13 A; 12 C; 3 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Recombinant swine pox virus - useful in vaccine for immunising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-130677/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cochran MD,
                                                                                                                  and to detect heartworm caused by D. immitis respectively.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Junker DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96US-0686968.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97WO-US12212
16.1%;
67.7%;
  Score 15; DB 19; Length 41; Pred. No. 5e+04;
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AAA37
                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                memorane fluidity; cold resistance; transgenic plant; baculovirus expression; PCR primer; ss.
                                                                                                                    Saccharomyces sp.
                                                                                                                                                                                                                                    Yeast acyltransferase YSCAT2 PCR primer, SEQ ID NO:190.
                                                                                                                                                                                                                                                                                                                                                          AAA37500 standard; DNA; 41 BP
                                                                             W0200018889-A2.
                                                                                                                                                                                             Acyltransferase; yeast; lipid synthesis; recombinant expression;
                                                                                                                                                                                                                                                                             15-AUG-2000 (first entry)
                                                                                                                                                                                                                                                                                                                   AAA37500;
24-SEP-1999;
                                      06-APR-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                         40 AATGTCTTGAATTTAAGATGATGGGTATGAG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33 AATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
  99WO-US22231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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25-SEP-1998;

98US-0101939

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to nucleic acids encoding novel plant C acyltransferase-like proteins (AAA37343-AA745) which comprise one of 8 CC conserved acyltransferase motifis (AAY99474-Y99481). Acyltransferases CC catalyse the transfer of acyl groups from a donor to a variety of c substrates such as glycerides, sterols, stanols and phosphatides. Such enzymes play a key role in lipid synthesis, and thereby affect the CC characteristics of the plant. For example, cold-hardened plants have compared to non-hardened plants, which makes the membrane more fluid and the plant CC non-hardened plants, which makes the membrane more fluid and the plant CC invention can be used as probes or for expressing acyltransferase-like CC proteins in host cells e.g., for recombinant protein production. They CC may be expressed in plant cells to alter the lipid composition of the CC the composition of plant oils. Sequences AAA77495-A37536 represent pCR capitansferase DNAs (AAA37527-A37534) for cioning into plant and capitaring any acceptance of the invention to amplify the yeast capitansferase DNAs (AAA37527-A37534) for cioning into plant and
                                                                                                                                                                                                                                                                                                                                                                                                AAD24539
                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
              27-JUN-1995;
                                         27-JUN-1996;
                                                                                              US6306649-B1
                                                                                                                                                                                                                                     Chimeric transcription factor; nuclear factor kappaB; NF-kappaB p65; biological material; biological research; biochemical purification; transgenic animal; gene therapy; acidic activation motif; ds.
                                                                     23-OCT-2001.
                                                                                                                                                                                                                    Unidentified.
                                                                                                                                                                                                                                                                                               12CA5 epitope DNA fragment #2.
                                                                                                                                                                                                                                                                                                                        07-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                             AAD24539 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                        AAD24539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 41 BP; 11 A; 10 C; 6 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 baculovirus expression constructs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 7; Page 32; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel acyltransferase related proteins useful for altering membrane fluidity in plant cells e.g. to induce chill tolerance .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-303447/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lassner MW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CALJ ) CALGENE LLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                             23 CCTGGGTGACAATGCCTTGGTTTCAACACTA 53
                                                                                                                                                                                                                                                                                                                                                                                                                                              7 CCTGCAGGTCAATGATTTTTTTTTCATCACAA 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Emig RA,
95US-000553P
95US-019614P
                                       96US-0672213.
                                                                                                                             /product= "12CA5 epitope fragment #2"
/note= "CDS does not include start and stop codon"
                                                                                                                                                               /*tag= a
                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16.1%;
67.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ruezinsky DM,

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15; DB 21
Pred. No. 5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Van Eenennaam A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 21; Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                           XX OS SOS XXX
                                                                                                                                                                                                                                                                                                                                                                                                                            밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
    WPI; 1997-402555/37.
                              Burkhardt M, Jiao J,
                                                                                                             30-JAN-1997;
                                                                                                                                         07-AUG-1997
                                                          (DADE-) DADE INT INC.
                                                                                    31-JAN-1996;
                                                                                                                                                                   WO9728191-A1
                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                            domo sapiens.
                                                                                                                                                                                                                                                                                                                                 AAT86970;
                                                                                                                                                                                                                                    tusion complex; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gilman MZ, Natesan S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P-PSDB; AAE15334.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002-033255/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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biological materials, in biological researches where precise control over a target gene is desired, such as in the expression of a protein or RNA of interest for biochemical purification and in tissue or organ specific expression of a protein or RNA in transgenic animals to elevate its biological function. It is also useful in gene therapy strategies by substantially elevating the expression of therapeutic target gene and transcription factors are useful for effecting transcription of target genes in genetically engineered cells or organisms containing them. The present sequence is 12CA5 epitope DNA fragment used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primer used in MHC fusion complex construction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a recombinant DNA sequence encoding a chimeric transcription factor. The transcription factor comprises one or more copies of a peptide sequence comprising all or part of a peptide sequence spanning 361-550 of human nuclear factor (NF)-kappaB p65 and a peptide sequence heterologous to the sequence selected from Vp16 V8, Vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp16 C, HSF or CTF. The recombinant DNA is used for producing vp16 B, Vp1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT86970 standard; DNA; 42 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 41 BP; 16 A; 6 C; 13 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New recombinant DNA sequence encoding a chimeric transcription factor, useful in gene therapy by elevating the expression of therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Column 31-32; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 target gene and allowing expression to reach therapeutically levels -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ARIA-) ARIAD GENE THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        45 TCAACACTATCGGAATGTGACTGTCGGGCCT 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 TCAACACTAGAGAGATGTGGCATGAAGGCCT 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               construction; major histocompatibility complex; MHC.
                                                                                                                                                                                                                                                                                                     96US-0596387
                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US01617.
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       Rhode PR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
Wong HC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 24; Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10;
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AAT7431
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SO CCC XXX PT PT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 42 BP; 6 A; 20 C; 11 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             arthritis, diabetes mellitus, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to an auto:immune disease, e.g. multiple sclerosis, rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single chain major histocompatibility complex comprising linked alpha and beta chains - useful for suppressing an immune response
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rapamyoin; responsive transfected cell; recombinant DNA; fusion protein; rapamyoin binding domain; RBD; FKBP; FKBP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Construct pCGNN-ZFHD1-1FRB fragment DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-OCT-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT74319;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Page 117; 217pp; English.
                                                                                  Clackson T, Gilman MZ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             misc_teature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                misc_teature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT74319 standard; DNA; 43 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is a primer used in the construction of major
P-PSDB; AAW23604
                                                                                                                                                                                                                                                                                           07-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                 WO9641865-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             histocompatibility complex (MHC) fusion complexes.
                            WPI; 1997-065453/06
                                                                                                                                               (ARIA-) ARIAD GENE THERAPEUTICS INC
                                                                                                                                                                                                     07-JUN-1995;
                                                                                                                                                                                                                              09-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    госат
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 16.1%; Score 15; Similarity 67.7%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SS
                                                                                                                                                                                                        95US-0481941
                                                                                                                                                                                                                                  96US-0598776.
                                                                                                                                                                                                                                                                                              96WO-US09948
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "FRB fragment that joins this sequence to the
3'-end of pCGNN-ZFHD1-1FRB fragment in AAT74320"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note- "ZFHD1 fragment that joins this sequence to the 5'-end of pCGNN-ZFHD1-1FRB fragment in AAT49028"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product= pcgnn-zfHD1-1fRB_fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'label- 5'-FRB_fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= c
/label= 3'-ZFHD1_fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      *tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 18; Length 42;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV03898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 second chimeric protein which is capable of forming a complex with rapamycin, or a rapamycin analogue, and the first chimeric protein and comprises at least one FKBP:rapamycin binding (FRB) domain and at least one domain heterologous to it. Contacting these genetically engineered cells with rapamycin or analogues results in the formation of a complex (between the fusion proteins, and the rapamycin), and initiation of a biological response. The products can be used for regulating biological events such as gene transcription, activation of an intracellular signal events such as gene transcription, activation of an intracellular signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence encodes the pCGNN-ZFHD1-1FRB construct fragment that contains the 3'-end of ZFHD1 and the 5'-end of FRB. The fragment is attached to AAT49028 by a missing fragment, and is attached to AAT4200 by another missing fragment which then forms the whole construct. This is used to produce an animal cell containing at least two recombinant DNAs. One DNA should encode a chimeric protein which is capable of binding to rapamycin, or a rapamycin analogue, and comprises at least one rapamycin binding domain (RBD) and at least one protein domain heterologous to it; the second recombinant DNA should encode a
                                                                                                                                                                                                                                                                                                                                                                                                     VZV; varicella-zoster virus; gE glycoprotein; epitope; peptide tag; chimeric protein; immunodetection; immunoprecipitation; glycoprotei antibody; PCR primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 43 BP; 16 A; 6 C; 13 G; 6 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transduction pathway leading to e.g. gene expression or apoptotic cell death, gene knock-out, blockade of a gene or inhibition of function of a gene product. They are used particularly for regulated gene therapy and for production of recombinant proteins and viruses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 63; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New rapamycin-responsive transfected cells - contg. recombinant DNA encoding fusion proteins which act as biological switches for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primer MP16 for chimeric protein construction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-APR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV03898;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV03898 standard; DNA; 45 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    regulating biological events
                                                                                                 Grose CF;
                                                                                                                                    (IOWA ) UNIV IOWA RES FOUND
                                                                                                                                                                              29-ЈUL-1996;
                                                                                                                                                                                                                      29-JUL-1996;
                                                                                                                                                                                                                                                            20-JAN-1998
                                                                                                                                                                                                                                                                                                      US5710248-A.
                                                                                                                                                                                                                                                                                                                                              Varicella-zoster virus
                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             45 TCAACACTATCGGAATGTGACTGTCGGGCCT 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6 TCAACACTAGAGAGATGTGGCATGAAGGCCT 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                 96US-0681935
                                                                                                                                                                                                                      96US-0681935.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.1%;
67.7%;

    Mismatches

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Pred. No. 5.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                            immunoprecipitation; glycoprotein gL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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Chimeric protein, useful for immunodetection and immunoprecipitation

WPI; 1998-109868/10.

comprising varicella zoster virus glycoprotein gE epitope linked

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                 contacting a solution containing a chimeric protein containing a protein linked to one or more heterologous peptide tags and which binds to the antibody to the peptide tag, contacting the antibody solution to the peptide tag, and then isolating the chimeric protein from the solution. A peptide tag, and then isolating the chimeric protein from the solution. A peptide tag, and then isolating the chimeric protein zoster virus (VZV) can be used as the peptide tag. The method and (I) are useful in the immunopurification and immunoisolation of proteins other than glycoprotein gE (the most abundant protein in the virion envelope). (I) may be used to identify cellular location of the chimeric protein in a cell, to identify the trafficking of the protein through cellular
                                                                                                                                                                                                                                                                                                 Immunoisolating and immunopurifying a protein comprises contacting a chimeric protein including a protein linked to one or more heterologous peptide tags with an antibody to the peptide tag \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents a primer used in the construction of the chimeric protein of the invention. This primer is used to amplify an antigenic fragment of the varicella-zoster virus glycoprotein gE. The chimeric protein is useful for immunodetection and immunoprecipitation, and comprises a protein linked to a heterologous peptide tag having the amino acid sequence shown in AAWAUGI, where the chimeric protein binds to antibody and to the peptide tag. Expressing a recombinant protein (especially VZV glycoprotein gL) as a chimeric protein facilitates its detertion and immunopurification
                                                                                                                                                                                                                     The invention relates to a method of isolating a protein that comprises
                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-432078/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (IOWA ) UNIV IOWA RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUL-2001
                                                                                                                                                                                                                                                                 Example 2; Column 15; 18pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6255462-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR mutagenic primer M16 rev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH46988 standard; DNA; 45 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH46988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 45 BP; 12 A; 9 C; 10 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Varicella-zoster virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection and immunopurification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Column 15; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .mmunoisolation; mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          госат
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30 GACAATGCCTTGGTTTCAACACTATCGGAATGTGACTGT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 GTCACCGTATTGCCTTTGACAGTATGCGATTGTGATAGT 45
to detect protein expression in transformed bacteria, as a marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       glycoprotein gE; VZV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96US-0681935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97us-0939323.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.1%; Score 15; 61.5%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide tag; immunopurification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.1e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 19; Length 45;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                 AAZ65554 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side
characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other
                                                                                                                                                                                                                                                                  Claim 3; Page 849; 2745pp; English.
                                                                                                                                                                                                                                                                                                            map of the human genome
                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-013267/01
                                                                                                                                                                                                                                                                                                                                                                                                                 Cohen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to quantify the level of gene expression from a cell, and to determine protein location within a cell. Antibodies to (I) can be used to test for the presence of the chimeric protein in a eukaryotic cell supernatant, and to identify, isolate, purify and quantify the amount of chimeric protein in a particular column fraction. Sequences AAH46983-991 represent primers used for incorporation of the 9E 3B3 epitope into a VZV protein gL by a recombination site specific PCR insertional mutagenesis method.
                                                                                                                                                                                                                                                                                                                             Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis; single nucleotide polymorphism; SNP; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ68554;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 45 BP; 12 A; 9 C; 10 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human map-related biallelic marker SEQ ID NO:2902.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-SEP-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30 GACAATGCCTTGGTTTCAACACTATCGGAATGTGACTGT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7 GTCACCGTATTGCCTTTGACAGTATGCGATTGTGATAGT 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                 Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0109732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-0082614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-IB00822
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.1%; Score 15; DB 22; Length 45; 61.5%; Pred. No. 5.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                 Chumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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        The invention relates to cold-adapted influenza viruses prepared by CC passage culture of A/X-31, B/Yamagata/16/88 or B/Lee/40 viruses at low CC temperatures. A cDNA gene of cold-adapted influenza virus HTCA-A101 can CC be selected from a group consisting of PB2 protein gene, PB1 protein CC gene (AAXB219-X82197). The method is useful for the production of cold-catively grown in fertilized eggs. M protein gene and NS protein CC adapted influenza virus that exhibit temperature sensitivity and can be catively grown in fertilized eggs. The virus is useful for vaccines for protection against 'flu. Live vaccines containing cold-adapted viruses CC have several advantages over killed vaccines. It can prevent reduction CC proteins would be denatured at its inactivation. It can also avoid CC proteins would be denatured at its inactivation. It can also avoid CC proteins of the vaccines the immunity by inducing 1gA and it can be caministered into a spray formulation via nasal cavity and thus its capplication is convenient for children. It is able to inhibit the CC growth of the wild-type virus and thus its therapeutic effect can be capplication sequences AAXB2198-X2221 represent RT-PCR primers used in the CC cloning of cold-adapted influenza virus HTCA-A101 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 91
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 47 BP; 11 A; 14 C; 7 G; 15 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cold-adapted influenza virus; passage culture; PB2 protein; PB1 protein; PA protein; NP protein; M protein; NS protein; temperature sensitivity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX82220;
                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 13; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    protective vaccines against flu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cold-adapted influenza viruses useful for the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-385377/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Youn Jw;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHEI-) CHEIL JEDANG CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vaccine; flu; influenza; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Influenza virus HTCA-Al01 gene cloning RT-PCR primer NS+.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-AUG-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX82220 standard; DNA; 48 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Influenza virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68 TCGGGCCTCCAGAAAACATTGAG 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13 TTGTGCCTCCATAAAATACTGAG 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kim HG, Kim J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97KR-0064854
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16.1%;
78.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kim SJ, Lee KH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seong BL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                 therapy. and in vaccine production. (1) and the polypeptides encoded by associated with inappropriate expression of polymorphic polypeptides.

CC chem may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides.

CC ror example, (1) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the polypeptides by expressing inactive proteins or to supplement the cc complementary sequences may also be used as DNA probes in diagnostic complementary sequences may also be used as DNA probes in diagnostic cassays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (1) may be used as antispens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity.

Crantipodies may also be used as diagnostic agents for detecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                     sequences (1), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAI75840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-NOV-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                          AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 902; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human silent SNP containing nucleic acid SEQ:2781.
presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-NOV-2000; 2000US-0726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-2000; 2000WO-US32758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 16.1%;
Local Similarity 67.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14 TCCATTCTGCCTGGGTGACAATGCCTTGGTT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 TCCATTATGTCTTTGTCACCCTGCTTTTGCT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0168138.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
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Query Match

Sequence 50 BP; 11 A; 11 C; 19 G; 9 T; 0 other;

Best Local

Similarity 67.

16.1%; Score 15; DB 22; 67.7%; Pred. No. 5.3e+04;

DB 22; Length 50;

Mismatches

0

Gaps

25 TGGGTGACAATGCCTTGGTTTCAACACTATC 55

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В
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                                                                                                                                                                                                           can determine a predisposition to cancers especially, prostate cancer.

Nucleic acid probes from the present invention are used to detect mismatches of the HPC1 gene or mRNA. Primers from the present invention can be used for determining the nucleotide sequence of a particular HPC1 allele. The mutations in the HPC1 allele which lead to an altered function of HPC1 are used in diagnostic and prognostic methods to determine a predisposition to cancer. HPC1 genes are also used in gene therapy to treat cancer. The HPC1 gene can be found in the HPC1 region of human chromosome 1, bound by the markers mM.GAAA.158;23.4 and comm.GA57e15.S16. AAA26513 to AAA26521 represent oligonucleotides used in hybrid selection in an example from the present incomplication. AAA2652 to AAA26537 represent PCR primers used in mutation screening of the HPC1 gene. AAA26538 to AAA26659 to AAA26652 encode the putative protein sequences given in AAA2657 to AAA26657 co AAA26657 to AAA26657 encode the putative protein sequences given in
                                                                                                                       Query Match
                                                                                                                                                          Sequence 19 BP; 3 A; 6 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a human prostate cancer susceptibility gene, designated the human prostate cancer predisposing gene 1 (HPC1). AAAJ2619 to AAAJ2610 represent nucleotide sequences from the HPC1 gene. The HPC1 gene sequences are used as diagnostic molecules since they
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel human prostate cancer predisposing gene in which somatic and germline mutations correlate with prostate cancer, useful in gene therapy and for diagnosis and prognosis of predisposition to prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; prostate cancer predisposing gene; HPC1; chromosome 1; susceptibility; tumour suppressor; prostate cancer; cytostati gene therapy; mutant; screening; PCR primer; ss.
                                                                                                                                                                                                   AAY82362 to AAY82365 from HPC1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; Page 73; 159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-270819/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tavtigian S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200012694-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human HPC1 mutation screening PCR primer SEQ ID NO:172.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA26628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA26628 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MYRI-) MYRIAD GENETICS INC
                                                                                                 POCAT
                                  16 CATTCTGCCTGGGTGACA 33
1 CAGCCTGCCTGGGTGACA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 TGGGTGCCCAGGCCAGGGTGTGAACTTTCTC 50
                                                                                                 Similarity
                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Teng DHF, Perry WL, Schroeder MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-0098183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US19508.
                                                                                                 88.98;
                                                                                                                15.9%;
                                                                              0
                                                                                             Score 14.8; DB 21;
Pred. No. 4.6e+04;
                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prostate cancer; cytostatic;
                                                                                                                Length 19;
                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Simard J;
                                                                           0;
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AAS17913/c
                                                                                                                                                                                Best Local Similarity
                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                 the heart failure is ischaemic or non-ischaemic and as a diagnostic market for bad clinical prognosis in heart failure or recurrent events after first event of heart failure. This sequence is the forward primer for real time PCR, labelled with SYBR green dye (not defined) and used with the reverse primer AASI/7912 to measure the quantity of IL-18 in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  manufacture of medicament for treatment and/or prevention of diseases or diseased conditions of the arteries, heart failure recurrent events, or as diagnostic marker for bad clinical prognosis in heart failure or recurrent events after first event of heart failure. The IL-18 inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  atherosclerosis; cardiant; vasotropic; antiulcer; myocardial infarction; atherosclerotic plaque; thrombosis of atherosclerotic plaque; stroke; ischaemic syndrome; heart failure; arteriosclerosis; vascularisation;
                                                                                                                                                                                                                                                                                             Sequence 21 BP; 5 A; 6 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prevention of atherosclerosis/arteriosclerosis; the manufacture of a medicament for treatment and/or prevention of thrombosis of atherosclerotic plaque (AP), AP ulcer, AP destabilisation (responsible atherosclerotic plaque (AP), AP ulcer, AP destabilisation (AP), AP ulcer, AP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treatment and/or prevention of atherosclerosis, thrombosis of atherosclerotic plaque, atherosclerotic plaque ulcer and heart failure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes the use of interleukin-18 (IL-18) inhibitor for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 28; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-MAY-2000; 2000EP-0109606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS17913 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                     invention.
                                                                                                                                                                                                                                                                                                                                                                                                     human atherosclerotic plaque samples, described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for stroke), atheroma, ischaemic syndromes e.g. myocardial infarction. AP disruption or heart failure recurrent events e.g. death, recurrent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                recurrent events
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of interleukin-18 inhibitor for manufacture of medicament for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-075227/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chvatchko Y, Tedgui A, Mallat Z;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISTF ) ARS APPLIED RES SYSTEMS HOLDING NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-APR-2001; 2001WO-EP04843.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200185201-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      atheroma; human; real time PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Interleukin 18; IL-18; cytokine; interleukin-18 inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS17913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              schaemia, re-vascularisation and progression of atherosclerosis, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      is useful for: the manufacture of a medicament for treatment and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Interleukin 18 (IL-18), forward primer for real time PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-MAR-2002 (first entry)
18 CTGGGAGACAATTCCTTG 1
                                                                    24 CTGGGTGACAATGCCTTG 41
                                                                                                                                            Conservative
                                                                                                                                                                        15.9%;
88.9%;
                                                                                                                                            0;
                                                                                                                                                                                                            Score 14.8; DB 24;
                                                                                                                                                                                Pred. No. 4.8e+04;
                                                                                                                                                Mismatches
                                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (responsible
                                                                                                                                            0,
                                                                                                                                            Gaps
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RESULT 95

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                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                              be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                               the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vascular endothelial growth factor receptor; VEGF receptor; flt-1; Vlascular endothelial growth factor receptor; receptor; flt-1; kDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms_like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 9; Page 140; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9715662-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX72318;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX72318 standard; RNA; 27 BP
Vascular endothelial growth factor receptor; VEGF receptor; flt-1;
                                 Human KDR VEGF receptor hammerhead ribozyme #581.
                                                                                                        AAX70909;
                                                                                                                                        AAX70909 standard; RNA; 27 BP
                                                                                                                                                                                                                                                                                                                                                             Sequence 27 BP; 9 A; 5 C; 4 G; 8 U; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes nucleic acid molecules which modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Escobedo J, McSwiggen J, Pavco P, Stinchcomb D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse flk-1 VEGF receptor hammerhead ribozyme #562.
                                                                   28-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                            27
                                                                                                                                                                                                                                                            61 GTGACTGTCGGGCCTCCAGAAAACATT 87
                                                                                                                                                                                                                                                                                               19;
                                                                                                                                                                                                                          GTGAGTGTTTCNTCATCAGAAAACATT 1
                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-0005974.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-0584040.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96WO-US17480.
                                                                                                                                                                                                                                                                                                            15.9%;

 Mismatches

                                                                                                                                                                                                                                                                                                          Score 14.8;
Pred. No. 5
                                                                                                                                                                                                                                                                                                            .2e+04;
                                                                                                                                                                                                                                                                                                                           DB 18;
                                                                                                                                                                                                                                                                                           8
                                                                                                                                                                                                                                                                                                                             Length 27;
                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                           Gaps
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X P X O O X
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WXXX
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                                                                                                                                                                                                                                                                                                                                  RESULT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AX67275 to AXX5752 represent specific examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 27 BP; 9 A; 5 C; 4 G; 8 U; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, % \left( 1\right) =\left( 1\right) ^{2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAY-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage;
tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                          Cellobiohydrolase; CBH A; CBH B; cellulose degradation; food production; condition: animal feed production; endoglucanase; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 9; Page 114; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-259017/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09715662-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                Primer for cellobiohydrolase coding sequence.
                                                                                                                                                                                                                      21-MAY-1999 (first entry)
                                                                                                                                                                                                                                                            AAX27056;
                                                                                                                                                                                                                                                                                            AAX27056 standard; DNA; 27 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes nucleic acid molecules which modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-JAN-1996;
26-OCT-1995;
                  WO9906574-A1
                                                      Aspergillus niger
                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Escobedo J, McSwiggen J, Pavco P, Stinchcomb D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CHIR ) CHIRON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ocal
                                                                                                                                                                                                                                                                                                                                                                                        27 GTGAGTGTTTCNTCATCAGAAAACATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                          61 GTGACTGTCGGGCCTCCAGAAAACATT 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                              19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15.9%;
Similarity 70.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96US-0584040
95US-0005974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 98
AAZ09713
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             04-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   conversion), wood pulp, paper and textiles. The cellulose in the material being treated may be degraded before and/or after ingestion. A particular use is in baking where addition of CBH to flour results in increased loaf volume and better texture (break and shred/crumb quality). The new CBH can be produced free of endoglucanases (which can affect fibre strength) so are better than crude cellulase compositions for fibre modification where strength retention is important (textile and paper manufacture and finishing). Cloned CBH and endoglucanases (from Mo9713862) should allow preparation of compositions tailored for particular applications. Unlike known CBH, the new enzymes have a broad ph optimum and are effective at the low ph values used in industrial processes, also they have better
                                         12-AUG-1999.
                                                                      WO9940117-A1
                                                                                                                                                     drug design; therapy; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; IgE; tissue damage; hypersensitivity; inflammatory cell recruitment; inflammatory modulator; Fc-gammaRIa; immune function regulation;
                                                                                                                                                                                             FC receptor; FC-gammaRIIa; human; FCR; model; three-dimension; 3-D; atomic coordinate; bloactive compound design; computer assisted;
                                                                                                                                                                                                                                          Human Fc-gammaRIIa PCR primer FI2.
                                                                                                                                                                                                                                                                                                                             AAZ09713 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 27 BP; 8 A; 10 C; 2 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cellobiohydrolase enzymes CBH A and CBH B of the invention.

The CBH proteins are used to degrade cellulose, optionally in combination with other enzymes. Particularly they are used in production, or processing, of foods, beverages (by liquefaction of plant cell wall material), animal feeds (particularly silage or grain-based feeds, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JUL-1998;
                                                                                                                Synthetic
                                                                                                                                         anti-inflammatory; immunoprotective; PCR primer; ss.
                                                                                                                                                                                                                                                                    15-NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents a PCR primer used to isolate DNA encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 5; Page 25; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New cellobiohydrolase enzymes from Aspergillus - used to degrade
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-153802/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               De Graaff LH, Gielkens MMC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-JUL-1997;
                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            thermal stability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               improve nutrient utilisation, and thus growth rate and/or feed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cellulose, e.g. in food, beverage, animal feed, pulp, paper and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KONN ) GIST-BROCADES BV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        textiles production or processing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Loca!
                                                                                                                                                                                                                                                                                                                                                                                                                           31 ACAATGCCTTGGTTTCAACACTATCG 56
                                                                                                                                                                                                                                                                                                                                                                                                    2 ACAATGCATTCCTTCCAAATCTACCG 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97EP-0202389
             99WO-IB00367
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15.9%;
73.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 20; Length 27; Pred. No. 5.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Visser J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                  YXX PPN XXX XXX PPN XXX XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
                                                                                                             30-DEC-1998.
                                                                                                                                                                                                                                                                                                 AAX06223;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Powell MS;
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hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIa and models of other FcR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to an Fc receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents a PCR primer used in the method of
Berno A, Chee M, Fan J, Lipshutz RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel model of an Fc receptor (FcR) protein representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of the FcR can be used to design bioactive chemical compounds, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphism; biallelic; paternity testing; forensic; genetic mapping; phenotypic typing; medicament; disease; marker; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human biallelic polymorphic DNA fragment TIGR-A003P30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotides, peptides, peptidomimetics and small organic molecules, by computer-assisted methods of drug design. Therapeutic compositions that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to
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                                                                                                                                                                                                Baell JB, Epa V, Garrett Maxwell KF, McCarthy TD,
                                                                                                                                                                                                                                        (AFFY-) AFFYMETRIX INC.
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06-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                        reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 313; 326pp; English.
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                                                                                                                                                                                                                                                                                                                                      Epa V, Garrett TPJ, Hogarth PM, Matthews BR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-0050594
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 Mismatches

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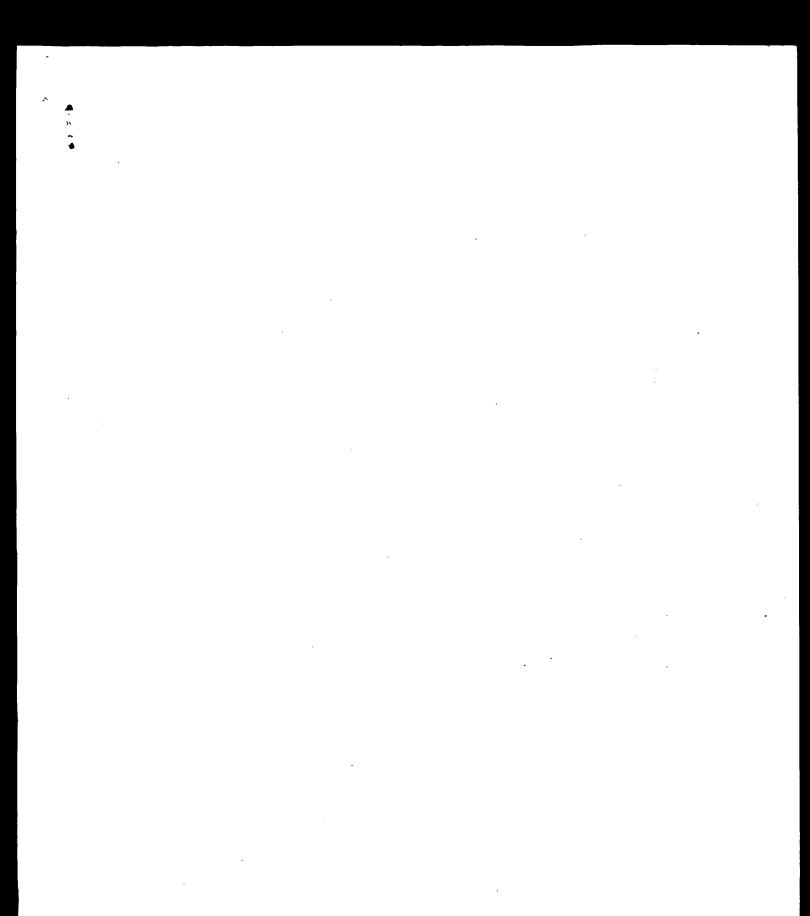
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             application programme and including records, each record comprising information identifying a polymorphism shown in the above sequences. The products and methods can be used for analysing polymorphic sites in individuals for testing for the presence of a disease phenotype or in forensics, paternity testing or genetic mapping of phenotypic traits. They can also be used for the production of polypeptides expressed by variant genes and for the production of transgenic animals. The nucleic acid segments can also be used in the manufacture of medicaments for the tratment or prohibits of alternative medical segments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            an individual. The invention further provides computer readable storage medium for storing data for access by an application programme being executed on a data processing system, such a method comprises a data structure stored in the computer readable storage medium, the data structure including information resident in a database used by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequences AAX06101-X06558 represent human DNA fragments which contain biallelic polymorphic markers. The base occupying the polymorphic site is indicated by the appropriate IUPAC-IUB ambiguity code. These fragments can be used in a method for determining polymorphic forms in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid segments containing polymorphic sites - used for, e.g. detecting a disease phenotype, in forensics, paternity testing or genetic mapping of phenotypic traits
Progestin-regulated gene - indicative of a subject's progestin responsiveness, also for treatment of metabolic disease or
                                                                 WPI; 1997-259026/23.
                                                                                                         Hamilton JA, Watts CKW;
                                                                                                                                                  (GARV-) GARVAN INST MEDICAL RES
                                                                                                                                                                                          19-JUL-1996;
24-OCT-1995;
                                                                                                                                                                                                                                                       24-OCT-1996;
                                                                                                                                                                                                                                                                                               01-MAY-1997.
                                                                                                                                                                                                                                                                                                                                          W09715674-A1
                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                      primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Progestin-regulated gene 1; PRG1; breast cancer; tumour; T-47D; hyperproliferation; metabolic disease; diabetes; cell cycle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Progestin regulated gene 1 (PRG1) reverse primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-JUL-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT60552;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT60552 standard; DNA; 36 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 15; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-080963/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                        antiproliferative; gene therapy; polymerase chain reaction; PCR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 15.9%; Local Similarity 67.9%;
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95AU-0006144.
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    Mismatches

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                                                                                                                                                                                             Search completed: July 5, 2003, 12:29:25
                                                                                                                         Matches
                                                                                                                                                         Query Match
                                                                                                                                                                                                                                      A forward primer (AAT60551) and reverse primer (AAT60552) were used to generate a full-length open reading frame of the novel human progestin-regulated gene PRGI (see also AAT60548) contg. terminal EcoRI and Sall restriction sites. The PCR product was cloned into bacterial expression vector pFLAG=AE-2 to allow prodn. of PRGI polypeptide (see also AAW15747) as a soluble fusion protein in
                                                                                                                                                                                         Sequence 36 BP; 7 A; 9 C; 10 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                   Example; Page 14; 50pp; English.
                                                                                                                                                                                                                             Escherichia coli DH5a transformants.
                                                                                                                                                                                                                                                                                                                                                                                                     hyperproliferation etc
                                                                                                                                    Match 15.9%;
Local Similarity 88.9%;
                                                                      73 CCTCCAGAAAACATTGAG 90
                                                   29 CCTCCAGGAAACACTGAG 12
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מח, 5.7e+04;
                                                                                                                                                   DB 18;
                                                                                                                       Indels
                                                                                                                                                     Length 36;
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                                                                                                                     Gaps
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Job time : 236 secs



Database: GenEmbl:* 1: gb_ba:* 2: gb_htg:* 3: gb_ln:* 4: gb_om:* 4: gb_om:* 5: gb_pat:* 6: gb_pat:* 6: gb_pat:* 6: gb_pat:* 7: gb_ph:* 9: gb_pr:* 10: gb_sy:* 11: gb_sts:* 11: gb_sts:* 12: gb_sy:* 13: gb_un:* 14: gb_vi:* 15: em_ba:* 16: em_fun:* 17: em_hum:* 19: em_ov:* 20: em_ov:* 21: em_ov:* 21: em_ov:* 22: em_ov:* 23: em_pat:* 24: em_ln:* 25: em_l:* 26: em_fu:* 27: em_sts:* 28: em_un:* 39: em_htg_ohum:* 31: em_htg_ohum:*	Post-processing: Minimum Match Maximum Match Listing first	Minimum DB seq length: 0 Maximum DB seq length: 50	Total number of hits satisfying	Searched: 2054640 seqs, 1	Scoring table: IDENTITY_NUC Gapop 10.0 , Ga	Title: US-09-843-377-3_COPY_10 Perfect score: 93 Sequence: 1 gagctgggagcactccattc.	Run on: July 5, 2003,	OM nucleic - nucleic search, us	Ge Copyright (c)
Im:* V:* Cher:* Lh:* 100% 1000% 1000 summaries		g chosen parameters: 841850	14551402878 residues	Gapext 1.0	3_COPY_1000_1092 ctccattcctccagaaaacattgaggtg 93	04:25:39 ; Search time 985 Seconds (without alignments) 2747.778 Million cell updates/sec	using sw model	GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.	

Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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                                                                                                                                                                     7 GGAGCACTCCATTCTGCCTGGG 28
                              Method for detecting tumor cells Patent: WO 0224745-A 4 28-MAR-2002;
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                      Abken, Hinrich (DE)
                                                                    Eukaryota;
                                                                           Homo sapiens
                                                                                                 AX404630.1 GI:21437911
                                                                                                          Sequence 4 from Patent WO0224745. AX404630
                                                                                                                        AX404630
                                                                                                                                                                                                                                                                  these organisms producing the same Patent: EP 0170204-A 33 05-FEB-1986; BOEHRINGER INSELHEIM INTERNATIONAL GMbH
                                                                                                                                                                                                                                                                                                                                                                                                              4 GCACTCCATCCAGCCTGGGCAACAA 28
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                                             Abken, H. and Schinkoethe, T.
                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
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Hauptmann, R., Meir
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                                                                                                                                                                                                 Similarity
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/organism="Homo sapiens"
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6 c 10 g
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                                                           Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 2.7e+04;
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                                                                                                                           Transgenic fish and vectors therefor Patent: US 5998697-A 14 07-DEC-1999;
                                                                                                                                                                                                                                                                     Sequence 14 from patent US 5998697 AR093240
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8 c 11 g
                                                           /organism="unknown"
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Accession number cg39667412"
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43 TTTCAACACTATCGGAATGTGACTGTCGGGC 73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (09-SEP-1997) Shuichi Asakawa, Keio University School of Medicine, Department of Molecular Biology; Shinanomachi 35, Shinjuka-ku, Tokyo 160, Japan (B-mail:asa@dmb.med.keio.ac.jp,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 (bases 1 to 34)
Asakawa, S. and Shimizu, N.
Direct Submission
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                                                                                                                                                                                                                                                               Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens STS NIB517, DH PROBE, FORWARD PRIMER, sequence tagged
                                                                                                                                                                             Pastan, I.H., Kreitman, R.J. and Puri, R.K.
Fusion proteins comprising circularly permuted ligands
Patent: US 5635599-A 31 03-JUN-1997;
                                                                                                                                                                                                                                            Unclassified
                                                                                                                                                                                                                                                                             Unknown.
                                                                                                                                                                                                                                                                                                            I44807.1 GI:2469520
                                                                                                                                                                                                                                                                                                                                          Sequence 31 from patent US 5635599
                                                                                                                                                                                                                                                                                                                                                            T44807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genomics 49 (2), 209-217 (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High-fidelity digital hybridization screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Asakawa,S. and Shimizu,N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens DNA.
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                                                  Similarity
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71.0%;
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                                                Score 16.6; DB 6; Length 39; Pred. No. 1.7e+05;
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35 TCTCAACACTCACCGGAGGTAACGGTGGGGC 5

PAT 08-AUG-2001

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                                                                                         50 bp
Sequence 5 from patent US 5639868.
147075
    1 (bases 1 to 50)
                  Unclassified
                                  Unknown
                                                Unknown.
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Janjic, N., Gold, L. and Tasset, D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unknown
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                                                                              I47075.1 GI:2471040
                                                                                                                                                                                                                                                                                                                                                                                     Patent: US 6177557-A 5 23-JAN-2001;
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AR125663
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                                                                                                                                                                                                                                                                                                                                                                                                              High affinity ligands of basic fibroblast growth factor and
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unclassified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unknown.
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d,L. and Beutel,B.
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Pred. No. 1.7e+05;
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Local Similarity 60.5%;
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Local Similarity 71.0%;
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                                          PR 23-OCT-1996 US 60/029592
PI WILLIAM V WILLIAMS, MICHAEL MADAIO, DAVID B WEHNER PC C12N15/09, A61K48/00//A61K39/00, C12N15/00
                                                                                                                                                                                                                                                                                                                                                                                                                                                          23;
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                                                                                                                                                                                                                                    unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               47 bp
Sequence 19 from patent US 6228621.
AR149818
                                                                                                                                             THE TRUSTEES OF THE UNIVERSITY OF PENNSYLVANIA
                                                                                                                                                                                                                                                               BD007169.1 GI:18635540 JP 2001503260-A/14.
                                                                                                                                                                                                                                                                                              Improved vaccines. BD007169
                                                                                                                                                                                         Williams, W.V., Madaio, M. and Wehner, D.B.
                                                                                                                                                                                                                                                     unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent: US 6228621-A 19 08-MAY-2001;
                                                                                                                                                          Improved vaccines
Patent: JP 2001503260-A 14 13-MAR-2001;
                                                                                                                                                                                                                                                                                                                          BD007169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unclassified
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                                                                        23-OCT-1997 JP 1998519728
23-OCT-1996 US 60/0295
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Location/Qualifiers
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PAT 31-JAN-2002

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23;
                                                      l (bases 1 to 40)
Chenchik,A., Zhu,Y., Diatchenko,L. and Siebert,P.
Methods and compositions for full-length cDNA Cloning using a template-switching oligonucleotide Patent: US 5962272-A 103 05-OCT-1999;
                                                                                                                                                                                                                                           Sequence 103 from patent US 5962272. AR078049
                                                                                                                                                                                                                                                                                                                                                                                                                                              26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  50 bp 1
Sequence 2480 from Patent WO0140521.
AX159152
                                                                                                                                                      Unclassified
                                                                                                                                                                                     Unknown.
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15 c 13 g 9 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Nucleotide deleted between bases 25 and 26
Accession number cg39667412"
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         /organism="unknown"
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/db_xref="taxon:9606"
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S74027
48 bp mRNA linear PRI 04-MAR-1:
TCR alpha VJ=islet-specific T-cell receptor alpha chain junction
region (clone K2.4) [human, type I diabetic patient, peripheral
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                                                                                                                              2 GTGCTGTGACCACTCAGTTTTCTGGTGGCTACAATAAGCTGATTT 46
                                                                                                                                                               1 GAGCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTT 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 GCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCT 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Durinovic-Bello, I., Steinle, A., Ziegler, A.G. and Schendel, D.J.
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1 (bases 1 to 48)
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/organism="Homo sapiens"
9 c 11 g 19 t
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/organism="Homo sapiens"
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/isolate="AS"
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/product="T-cell receptor alpha-chain joining region"
/protein_id="CAA82200.1"
/db_xref="GI:607119"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /cell_type="CD4+ T-cell clone"
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 17.4%; Score 16.2; DB 9; Local Similarity 60.0%; Pred. No. 2.5e+05;
                                                                                             Curagen Corporation (US)
                                                                                                             methods of use thereof Patent: WO 0151670-A 532 19-JUL-2001;
                                                                                                                                                                                                                                                                                                                               Sequence 532 from Patent W00151670
                                                                                                                                                                                                                                                                                                                                                                                                                                                  N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GAGCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTT 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GenBank staff at the National Library of Medicine created this entry [NCBI gibbsq 157354] from the original journal article. This sequence comes from Fig. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 48)
Durinovic-Bello,I., Steinle,A., Ziegler,A.G. and Schendel,D.J
HLA-DQ-restricted, islet-specific T-cell clones of a type I
diabetic patient. T-cell receptor sequence similarities to
insulitis-inducing T-cells of nonobese diabetic mice
Diabetes 43 (11), 1318-1325 (1994)
                                                                                                                                              Shimkets, R.A. and Leach, M.D.
Nucleic acids containing single nucleotide polymorphisms and
                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   blood lymphocytes, mRNA Partial, 48 nt].
S74027
                                                                                                                                                                                      Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
1 (bases 1 to 50)
                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                   AX199602.1 GI:15390036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens peripheral blood lymphocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                GTGCTGTGACCACTCAGTTTTCTGGTGGCTACAATAAGCTGATTT 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
/organism="Homo sapiens"
/db_xref="taxon:9606"
25. .26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /translation="CAVTTQFSGGYNKLIF"
                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /product="islet-specific T-cell receptor alpha chain
junction region"
/protein_id="AAB32155.1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="This sequence comes from Fig. 3"
/codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /partial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="islet-specific T-cell receptor alpha chain junction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:9606"
/note="type: I diabetic patient"
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                                                           SOURCE
                                                                           KEYWORDS
                                                                                                                                DEFINITION
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                                                                                                                                                                                      RESULT 19
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                                                                                                                 ACCESSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      source
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                                                                                                                                                                                                                                                                                                                   28;
                                         Unknown
                                                           Unknown.
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 TGCCTGGGTGACAATGCCTTGGTTTCAACACT 52
                                                                                                                                                                                                                                                                                             49 AGCTTGAGCCACCGCACCAGGCCAGGATCATAATTTTTATTTGCAAC 2
Unclassified.
1 (bases 1 to 40)
                                                                                                                                                                                                                                                                                                                                      2 AGCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAAC 49
                                                                                                                                 Sequence 6 from patent US 6180391.
AR127202
                                                                                                                AR127202.1 GI:14113795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence of the flanking viral DNA was: 5-TCTTGTCTTCGTTGGGAGTGAATTAGCCCTTCCA-3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                genomic DNA was cut by HindIII; the viral-cellular junctions were amplified by using the inverse PCR method; the first five bases (1-5) are duplicated during the HIV DNA integration process; the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (22-JAN-1998) Infectious Disease Laboratory, The Salk Institute for Biological Studies, 10010 North Torrey Pines Rd, La
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 50)
Carteau, S., Hoffmann, C. and Bushman, F.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chromosome structure and HIV-1 cDNA integration: centromeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens clone SUPTH34 sequence flanking the HIV-1 provirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jolla, CA 92037, USA
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Carteau,S., Hoffmann,C. and Bushman,F.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone="SUPTH34"
/cell_line="supT1"
/cell_q c 13 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
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18 c     14 g     5 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                      17.2%;
58.3%;
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                                                                                                                                                                                                                                                                                                                                                                   Score 16;
Pred. No.
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                                                                                                                                                                                 linear
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                                                                                                                                                                                 PAT 16-MAY-2001
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BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11 CACTCCATTCTGCCTGGGTGACAATGC 37
                                                                                                                                                                                                                                                                                                                                          32
                                                                                                                                                                                                                                                                                                                                                                43 TTTCAACACTATCGGAATGTGACTGTC 69
43 TTTCAACACTATCGGAATGTGACTGTC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27 CACTCAGTTCGTACTGGTTGACAAGGC 1
                        20; Conservative
                                                                                                                                    1 (bases 1 to 42)
Lemieux,B., Landry,B.S. and Sapolsky,R.J.
Brassica polymorphisms
Patent: US 6358686-A 45 19-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                            20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brassica polymorphisms
Patent: US 6114116-A 45 05-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent: US 6180391-A 6 30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brown, W.C.
                                                                                                                                                                                                                                                   Sequence 45 from patent US 6358686 AR200741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 45 from patent US 6114116. AR109086
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                                                                                                                                                                                                                                       AR200741.1 GI:20251629
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Lemieux, B., Landry, B.S. and Sapólsky, R.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unknown
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                                                                                                                                                                                                     Unknown
                                  h 17.0%; Score 15.8; DB 6; Length 42; Similarity 74.1%; Pred. No. 3.5e+05;
                                                                                                                                                                                                                Unknown.
                                                                                                                                                                                                                                                                             AR200741
                                                                                                                                                                                                                                                                                                                                                                                                     17.0%;
Similarity 74.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Inclassified.
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                                                                                                /organism="unknown"
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74.18;
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                       0; Mismatches

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Pred. No. 3.
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AX417271
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Harkki, A.M., Myasnikov, A.N., Apajalahti, J.H.A. and Pastinen, O.A.
Manufacturing of xylitol using recombinant microbial hosts
Patent: US 5631150-A 8 20-MAY-1997;
Location/Qualifiers
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                                                        Sequence 14 from patent US 5792456. AR022050
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Unknown
                                    AR022050.1 GI:3976112
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OS Unidentified
PN JP 1999196888-A/1
PD 27-JUL-1999
PF 16-JAN-1998 JP 19
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Yelton,D., Glaser,S., Huse,W. and Rosok,M.Joanne.
Yelton,D., Glaser,S., Huse,W. and Rosok,M.Joanne.
Mutant BR66 antibodies reactive with human carcinomas
Patent: US 5792456-A 14 11-AUG-1998;
Location/Qualifiers
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                                                                                            Patent: JP 1999196888-A 10 27-JUL-1999;
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JP 1999196888-A/10.
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                                                                                                                                    Takashi, K., Makoto, G., Masato, T. and Hideaki, Y.
                                                                                                                                                                                unidentified
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                                                                                                                      Process for producing organic acid by pyruvate carboxylase gene
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                                                                                                                                                  (bases 1 to 35)
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TAKASHI KOBAYASHI, MAKOTO GOTO, MASATO TERASAWA, HIDEAKI YUKAWA
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Pred. No. 4
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y Match 16.8%;
Local Similarity 81.8%;
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Sequence 4 from Patent WO0246456.
AX457962
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AX457962.1 GI:21724857
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Ramchandani,S., Bigey,P. and Szyf,M.
Genomic structure of the human DNA methyltransferase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          H3G-1Y6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (08-NOV-1999) Pharmacology and Therapeutics, McGill
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens DNA methyltransferase (DNMT1) gene, intron 9.
AF202558
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CC Topology: Linear
                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University, 3655 Promanade Sir William Osler, Montreal, Quebec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ramchandani, S., Bigey, P. and Szyf, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                          TTCTGTTTGGGTGTTGGTTTGGTTTGA 37
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                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C12P7/40,C12N1/21,C12N15/09//(C12P7/40,C12R1:13),(C12N1/21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
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                                                                                                                                                                                                                                                                                                                                             /note="contains intron 9"
6 c 13 g 22 t
                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                  /gene-"DNMT1"
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9 c 7 g
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                                                                                                                                                                                                                                                          16.8%;
70.0%;
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 Mismatches

                                                                                                                                                                                                                                                             Score 15.6;
Pred. No. 4
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                                                                                                                                                                                                                                                                                  Length 43;
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                 SOURCE
                              KEYWORDS
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                                                                                                                         RESULT 30
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TITLE
ORGANISM
                                                                                                                                                                                                                                     Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                misc_feature
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                                                                                                                                                                                                 16 CATTCTGCCTGGGTGACAATGCCTTGGTTT 45
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                                                                                                                                                                    12 CCTGGTGCCTGGGCCACCATGCCATGCTCT 41
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                                                                                         A46551
unidentified
                                                            Sequence 7 from Patent W09524487.
             unidentified.
                                            A46551.1 GI:2300716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Curagen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimkets, R.A. and Leach, M. Nucleic acids containing single nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 612 from Patent WO0140521. AX157284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isothermal amplification of nucleic acids on a solid support Patent: WO 0246456-A 4 13-JUN-2002; Applied Research Systems ARS Holding N.V. (AN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct. synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AX157284.1 GI:14538615
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                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO 0140521-A 612 07-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of use thereof
                                                                                                                                                                                                                                                                                                            /note="2 of 2 allelic variants (611 is other entry)" 16 c 17 g 9 t
                                                                                                                                                                                                                                                                                                                                                              Accession number cg20611295"
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7 c 17 q 4 t
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                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="taxon:32630"
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                                                                                       linear
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                                   AUTHORS
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                                                                                                                                                                                                                                                                                 34 ATGCCTTGGTTTCAACACTATCGGA 58
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                                                                                                                                                                                                                                                                                                                               19;
Gold,L., Janjic,N. and Pagratis,N. High affinity oligonucleotide ligands to growth factors Patent: US 6207816-A 285 27-MAR-2001;
                                                                                                                                               Sequence 285 from patent US 6207816. AR141007
                                                                                                                                                                                                                                                             7 ATCCCTTGGTTGCAACAGCAGCTGA 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent: WO 9524487-A 7·14-SEP-1995;
HOECHEY SCHERING AGREVO GMBH (DE)
Other publication DE 443821 960425
Other publication DE 4435366 960328
Other publication AU 2067995 950925
Other publication DE 4408629 950914.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 31)
Mueller-Roeber, B., Landschuetze, V. and La, C.U.
PROCESSES FOR INHIBITING AND FOR INDUCING FLOWER FORMATION IN
                                                                                 Unknown
                                                                                                                               AR141007.1 GI:14483503
                                                                                                                                                                               AR141007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent: WO 9806831-A 6 19-FEB-1998; MAX PLANCK GESELLSCHAFT (DE); WILLMITZER LOTHAR (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            La,C.U. and Willmitzer,L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unidentified
                                                                 Unclassified
                                                                                               Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TRANSGENIC PLANT CELLS AND PLANTS WITH MODIFIED ACETYL-COA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            unidentified
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                                               (bases 1 to 41)
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/db_xref="taxon:32644"
8 c 9 g
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                                                                                                                                                                                                                                                                                                                                           16.6%;
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                                                                                                                                                                                                                                                                                                                                                                                              Chen H.M., Rossler C., Chrast R., Antonarakis S.E.; "Cloning of 559 potential exons of genes of human chromosome 21 by exon trapping.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (17-MAR-1995) to the EMBL/GenBank/DDBJ databases. Stylianos E. Antonarakis, Division of Medical Genetics, University and Cantonal Hospital of Geneva, CMU, 1 rue Michel-Servet, 1211 Geneva,
                                                                                                                                              Sequence 43 BP; 8 A; 14 C; 9 G; 7 T; 5 other;
                                                                                                                                                                                                              exon
                                                                                                                                                                                                                                                                                              source
                                                                                                                                                                                                                                                                                                                                    Key
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chen H.M., Rossier C., Chrast R., Antonarakis S.E.; "Cloning of trapped exons from human chromosome 21";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           trapped exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   H.sapiens DNA for trapped exon (ID HMC37D09)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-JUL-1996 (Rel. 48, Created)
29-MAY-1997 (Rel. 52, Last updated, Version 3)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         X88097.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antonarakis S.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (human)
                                                       Local Similarity 57.9%; Pred. No. 5.1e+05; ses 22; Conservative n. Minner.
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                   55 CGGAATGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
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                                                                                                                                                                                                                              /organism="Homo sapiens"
                                                                                                                                                                                   /note="trapped exon"
                                                                                                                                                                                                                                                    /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                        /chromosome="21"
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Local Similarity 71.4%;
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Local Similarity 66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22 GCCTGGGTGACAATGCCTTGGTTTCAACACTAT 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43 CTGAGAGTGACTCGGGAGNCTNCAGANTNNAGCGTGGT 6
AR051493 31 bp
Sequence 63 from patent US 5830670.
                                                                                                     3 GATGTACGAGGACTCCATCCTGGCTGTG 30
                                                                                                                                   1 GAGCTGGGAGCACTCCATTCTGCCTGGG 28
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Sequence 27 from Patent W00109303
AX080977
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                                                                                                                                                                                                                                                                                                                                                                                          Hauptmann,R., Meindl,P., Dworkin-Rastl,E., Adolf,G.D., Swetly,P.,
Pieler,C. and Hauel,N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleotide sequence 34 from patent number EP0170204.
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RESULT 38
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               Patent: US 5948888-A 63 07-SEP-1999;
                                                      1 (bases 1 to 31) de la Monte, S. and Wands, J.R.
                                                                                                                                                  Sequence 63 from patent US 5948888 AR073178
                                        Neural thread protein gene expression and detection of Alzheimer's
                                                                                                 Unknown
                                                                                                                                     AR073178.1 GI:9999941
                                                                                                                                                                                                                                                                                                                                                                                                                 Patent: US 5948634-A 63 07-SEP-1999;
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                                                                                 Unclassified.
                                                                                                            Unknown.
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de la Monte, S. and
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Pred. No. 5.8e+05;
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y Match 16.1%; Score 15; DB 6; I Local Similarity 100.0%; Pred. No. 6.2e+05; hes 15; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                        Chesnut, J.D., Shuman, S., Madden, K.R., Heyman, J. and Bennett, R.P. Methods and reagents for molecular cloning Patent: WO 0216594-A 15 28-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 from Patent W00216594. AX468447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human single nucleotide polymorphisms
Patent: WO 0166800-A 694 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US)
                                                                                                                                                                                                                                                                                                                                                                                synthetic construct
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10 c 8 g
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/db_xref="taxon:32630"
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85.0%;
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PAT 03-APR-1996

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                Sequence 7 from patent US 5981262.
                                                                                                                                                                                                                                                                                                                               Human single nucleotide polymorphisms
Patent: WO 0166800-A 1732 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US)
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   AR085045.1 GI:10011816
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Sequence 8 from patent US 5476926.
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AR127205
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Spiegelman, B.M., Graves, R. and Ross, S. Adipocyte-specific DNA sequences and use thereof in the production of transgenic animals exhibiting altered fat tissue metabolism patent: US 5476926-A 12 19-DEC-1995;
                                                                                                       Brown, W.C
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                                                             Patent: US 6180391-A 9 30-JAN-2001;
                                                                                                                                    Unclassified
                                                                                                                                                    Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brugge, J., Morganstern, J., Shiue, L., Zydowsky, L., Zoller, M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unclassified.
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                                                                                       Highly efficient controlled expression of exogenous genes
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ARI11609
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Cochran, M.D. and Junker, D.E.
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                                                                                                                                                                       Sequence 143 from patent US 6251403. AR158235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unknown.
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                                         Recombinant swinepox virus Patent: US 6251403-A 143 26-JUN-2001;
                                                                    Cochran, M.D. and Junker, D.E.
                                                                                                 Unclassified.
                                                                                                                 Unknown
                                                                                                                               Unknown
                                                                                                                                                         ARI58235.1 GI:16220250
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Patent: US 6221361-A 70 24-APR-2001;
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     AR175028
                                                                     1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31
                                                                                                                            21; Conservative
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Rhode, P. R., Jiao, J. -A., Burkhardt, M. and Wong, H.C.
Single chain MHC complexes and uses thereof
Patent: US 5869270-A 38 09-FEB-1999;
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Sequence 38 from patent US 5869270.
AR033895
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Patent: US 6306649-A 30 23-OCT-2001;
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Gilman, M.Z. and Natesan, S.
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                                                                                                                                                                                                                                                                                                           Unclassified.
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Local Similarity 67.7%;
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          Peptide tag for immunodetection and immunopurification Patent: US 6255462-A 13 03-JUL-2001;
                                                                                                                                                                                                                                 1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31
                                                                                                                                                                                                                                                                                                                                                                                                                          Chavaillaz, P.A., Edwards, A.C., Gramme Weidanz, J.A. and Wong, H.C. Mccomplexes and uses thereof Patent: EP 0997477-A 38 03-MAY-2000; SUNOL MOLECULAR CORP (US)
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Rhode.P.R., Jiao, J.-A., Burkhardt,M. and Wong,H.C.
MHC molecules and uses thereof
Patent: US 6309645-A 38 30-OCT-2001;
                                         Grose, C.F
                                                                                                                       AR161438.1 GI:16227322
                                                                                                                                                 Sequence 13 from patent US 6255462.
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Sequence 38 from Patent EP0997477.
AX032460
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67.7%;
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Pred. No. 7.3e+05
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l Similarity 61.5%;
24; Conservative
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Sequence 2781 from Patent WO0140521.
AX159453
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I81454
                                                                                                                                                                                                                    Patent:
                                                                                                                                                                                                                                          Nucleic acids containing single nucleotide polymorphisms and
                                                                                                                                                                                                                                                                 Shimkets, R.A. and Leach, M.
                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 61.5%;
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                                           /note="1 of 2 allelic variants (2782 is other entry)" 11 c 19 g 9 t
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                                                                                        /note="Nucleotide deleted between bases 25 and 26
/ccession number cg42462239"
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Pred. No. 7.3e+05;
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                                 61 GTGACTGTCGGGCCTCCAGAAAACATT 87
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27 GTGAGTGTTTCNTCATCAGAAAACATT 1
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                                                                19; Conservative
                                                                                                                                                                                                                    1 (bases 1 to 27)
Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 3659 12-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                        27 bp Dt
Sequence 3659 from patent US 6346398.
AR188171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 8 from Patent WO0185201.
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/db_xref="taxon:9606"
a 6 c 5 g !
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88.9%;
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   and Hulett, M.D.
                1 (bases 1 to 30)
McKenzie, I.F.C., Hogarth, M.P., Hibbs, M.L., Scott, B.M., Bonadonna, L.
                                                            Unknown
                                                                                                                        Sequence 33 from patent US 5985599 AR086544
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Sequence 5068 from patent US 6346398.
AR189580
                                               Unclassified
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CELLULOSE DEGRADING ENZYMES OF ASPERGILLUS
PATENT: W 9906574-A 11 11-FEB-1999;
VISSER JACOB (NL); GIELKENS MARCUS MATHEUS CATHAR (NL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unidentified
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                                                                             Unknown.
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Pred. No. 8.1e+05;
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Pred. No. 8.1e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                     73 CCTCCAGAAAACATTGAG 90
                                                                                                                                 Ehrlich, R., Rotem-Yehudar, R. and Laham, N.
A soluble beta 2 microglobulin (beta2m)/hfe monochain for biotechnological and therapeutic applications Patent: WO 0224929-A 12 28-MAR-2002;
                                                                                                                                                                                                                          synthetic construct
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Patent: EP 0865495-A 5 23-SEP-1998;
GARVAN INST MED RES (AU)
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                                                                                                                                                                                                                                       synthetic construct.
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                                                                                                                                                                                                                                                                                                  Sequence 12 from Patent W00224929.
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                                                                                                                                                                                                         artificial sequences.
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/note="synthetic"
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                                                                                                                                                                              1 (bases 1 to 41)
Lemieux,B., Landry,B.S. and Sapolsky,R.J.
Brassica polymorphisms
Patent: US 6358686-A 147 19-MAR-2002;
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   6 GTGAAAAGGCATTGTKCTCCAAAATCTCGCTCTGCG 41
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Patent: WO 0159103-A 4860 16-AUG-2001;
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Sequence 147 from patent US 6358686.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RIBOZYME PHARMACEUTICALS, INC. (US) ; Blatt, Lawrence (US) ; McSwiggen, James (US) ; Chowrira, Bharat M. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blatt,L., Mcswiggen,J. and Chowrira,B.M. Method and reagent for the modulation and diagnosis of cd20 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AX219418.1 GI:15547142
                                                                   h 15.9%; Score 14.8; DB 6; Similarity 61.1%; Pred. No. 8.7e+05;
                                                                                                                                                                                                                                        Unclassified
                                                                                                                                                                                                                                                                   Unknown.
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Similarity 62.9%;
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31
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Pred. No. 8.6e+05;
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TITLE
Query Match
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                                                                                                                                                                                                                                                                                                         TITLE
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                                                                                 source
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Match 15.9%;
Local Similarity 64.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CAACACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic DNA related to mink growth hormone E06872
                                                                                                                                                                                                                                                                                                                                                                                                                                             FRAGMENT FOR GENE SECRETION MANIFESTATION Patent: JP 1994070776-A 1 15-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Harada,Y., Nakano,E., Tatsumi,H. and Umetsu,M.
NEW RECOMBINANT DNA, PRODUCTION OF MINK GROWTH HORMONE AND DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     artificial sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               synthetic construct.
synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JP 1994070776-A/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Curagen Corporation (US)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 478 from Patent W00151670.
                                                                                                                                                                                                                                                                                                                                                                                                                          KIKKOMAN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent: WO 0151670-A 478 19-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                    HARADA YASUHIRO, NAKANO EIICHI, TATSUMI HIROKI, UMETSU MOTOAKI C12N15/18,C12P21/02,(C12P21/02,C12R1:865); strandedness: Single;
                                                                                                                                                                            anti-sense: No.
                                                                                                                                                                                                                                                                                                                                                       JP 1994070776-A/1
                                                                                                                                                                                                                                                                                                                                                                          Artificial gene
Artificial sequence; Genes.
                                                                                                                                                                                                                                                                                                            26-AUG-1992 JP 1992227507
                                                                                                                                                                                                  hypothetical: No;
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a
                                                                                                                                                                                                                     copology: Linear;
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Accession number cg43321506"
a 13 c 17 g 8 t
                                                                                    /db_xref="taxon:32630"
                                                                                                                                                  Location/Qualifiers
                                                                                                         organism="synthetic construct"
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15.98;
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Pred. No. 8.8e+05;
Score 14.8;
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DB 6;
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Length 45;
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VERSION
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MEDLINE
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                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                    51 CTATCGGAATGTGACTGTCGGGCCTCCAGAAAAC 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19 TCCAGCCTGTTTGCCAATGCCGTGCT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 TTCTGCCTGGGTGACAATGCCTTGGT 43
      Homo sapiens
                                                                                                                                                                                                              2 CTCTTGGGACGGGGGTGAGGGGACTCCAGGACAC 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     שניטיטים 45 bp mRNA linear PRI 03-OCT-
H.sapiens (patient JoJ, clone 2) mRNA for T-cell receptor delta
chain V-J region.
                                                              A69934.1 GI:4774438
                                                                                                  Sequence 29 from Patent WO9807850
                                                                                                                      A69934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RELATED SEQUENCES: Y00793; P.N.A.S, 85:6097-6101 (1988); P.N.A.S, 85:9714-9718 (1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Howard,R.F. and White,B.
Expansion of selected V delta 1+ gamma delta T cells in systemic sclerosis patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                            human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 (bases 1 to 45)
Yurovsky, V.V., Sut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (21-MAY-1993) V.V. Yurovsky, University of Maryland at Baltimore, 10 South Pine Street, MSTF, Rm 8-22, Baltimore, MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         X72133.1 GI:556771
                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       joining region; T cell receptor; T-cell receptor delta chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunol. 153 (2), 881-891 (1994)
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                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                 /product="T cell receptor delta chain V-J region" 11 	ext{ c} 	ext{ 17 g} 	ext{ 7 t}
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /gene="Vdeltal-Jdeltal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /cell_type="gamma /delta T cell"
/tissue_type="blood"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                     /gene="Vdeltal-Jdeltal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="rearranged"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /chromosome="14"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                      15.9%;
64.7%;
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                                                                                                                                                                                                                                                                                                      Score 14.8; DB 9; Length 45; Pred. No. 8.8e+05;
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                                                                                                                    PAT 07-MAY-1999
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                                                                                                                                                                                                                                RESULT 72
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 47)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14 TCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAACACTATC 55
                                                                                                                                                                                                                                                                                      9
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Sequence 492 from Patent W00151659.
AX195022
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Preddie, E.R. and Bergmann, J.E.
                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                        9 AGCACTCCATTCTGCCTGGGTGACAATG 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 TCCTTTCTTTCAGAATGCCTATTCCTGTGCATTAAAAGTGTC 42
                                                                                              JP 2001503968-A/21.
                                                                                                                                                                                          BD010500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENSET (FR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     involved in central nervous system disorders
Patent: WO 0151659-A 492 19-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chu, T., Blumenfeld, M. and Cohen, D.
                                                                                                                    BD010500.1 GI:18638873
                                                                                                                                                  Alzheimer's disease and Down syndrome in humans
                                                                                                                                                                        Agents for pre-symptomatic detection and therapeutic targeting of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Biallelic markers derived from genomic regions carrying genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AX195022.1 GI:15385669
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGENTS FOR PRE-SYMPTOMATIC DETECTION AND THERAPEUTIC TARGETING OF
                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                      ACCAGTGCATTTTGCMTCTGAGACATTG 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="18-867-331 : polymorphic base A or C"
12 c 8 g 18 t 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:9606"
/clone="PALZ11"
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                                              13 ATTGGACTGTGAGTCTGGTGCCCTAAGCTAACGT 46
                                                                                53 ATCGGAATGTGACTGTCGGGCCTCCAGAAAACAT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14 TCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAACACTATC 55
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PC A61K3
CC Strar
CC Topol
FH Key
FT Sourc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 TCCTTTCTTCAGAATGCCTATTCCTGTGCATTAAAAGTGTC 42
                                                                                                                                                                                                                                                                                                                                                                Chuba, P.J., Pelz, K., Krekeler, G., de Isele, T.S. and Gobel, U. Synthetic oligodeoxynucleotide probes for the rapid detection of bacteria associated with human periodontitis.

J. Gen. Microbiol. 134 (Pt 7), 1931-1938 (1988)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Actinobacillus hominis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A.hominis 16S rRNA, V8 variable region.
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                                                                                                                                                                                                                                                                                                                                       2469776
                                                                                                                                                                                                                                                                                                                                                                                                                                         Actinobacillus.

1 (bases 1 to 49)
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                                                                                                                                        Similarity
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Similarity 59.5%;
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                                                                                                                       Conservative
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C12N15/12, C12N15/55, C12N9/64, C07K14/47, C12Q1/68, G01N33/50,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism='Homo sapiens (human)'.
Location/Qualifiers
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22-AUG-1996 CA 21839
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JP 2001503968-A/21
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                                                                                                                                                                                                            /note="16S rRNA, V8 region"
8 c 17 g 15 t
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/db_xref="taxon:9606"
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64.7%;
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Pred. No. 8.9e+05;
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Pred. No. 8.9e+05;
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AUTHORS
TITLE
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AX157274
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                                                                                                                                                                                                        Curagen
                                                                                                                                                                                                                         Patent:
                                                                                                                                                                                                                                         methods
                                                                                                                                                                                                                                                                         Shimkets, R.A. and Leach, M.
                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                     AX157274.1 GI:14538605
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 602 from Patent W00140521. AX157274
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AX157274
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (26-JAN-1999) Agroecology, University of British Columbia, 187 - 2357 Main Mall, Vancouver, BC V6T 124, Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 (bases 1 to 50)
Pang, S.W.Y., Cheng, K.M. and Ritland, C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pang,S.W., Ritland,C., Carlson,J.E. and Cheng,K.M.
Japanese quail microsatellite loci amplified with chicken-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae; Coturnix.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Coturnix japonica. Coturnix japonica
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                                                                                                                                                                                                                                                    Nucleic acids containing single nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
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99371928
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                                                                                                                                                                                                        Corporation (US)
                                                                                                                                                                                                                   of use thereof WO 0140521-A 602 07-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="synonym:Coturnix japonica"
11. .27
                                                                         /note="Nucleotide deleted between bases 25 and 26
Accession number cg6586279"
                                      /note="2 of 2 allelic variants (601 is other entry)"
                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Coturnix japonica"
/db_xref="taxon:93934"
                                                                                                                               /db_xref="taxon:9606"
                                                                                                                                                  organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="microsatellite UBC0006"
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Indels

0; Gaps

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linear

PAT 22-JUN-2001

Query Match

15.9%;

Score 14.8;

DB

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Length 50;

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RESULT 78
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                                                                   71 GGCCTCCAGAAAACATTGAGG 91
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3730 from Patent W00129262
AX118607
                                                                                                                                                                                                                                                     Genotyping reagents, kits and methods of use thereof patent: WO 0129262-A 3730 26-APR-2001; Orchid BioSciences, Inc. (US)
                                                                                                                                                                                                                                                                                                                                         synthetic construct
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LUNA SUSANA DE (ES); THANGUE NICHOLAS BARRIE (GB)
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De.L.L. and La.T.N.
Dep transcription factor-interacting protein and its use
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Picoult-Newburg, L. and Pohl, M.
                                                                                                                                                                                                                                                                                                                        artificial sequences.
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BOEHRINGER INGELHEIM INTERNATIONAL GmbH
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 unidentified
                            Al1501.1 GI:492485
                                                         Nucleotide sequence 41 from patent number EP0170204
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Pieler, C. and Hauel, N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unidentified.
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               Materials and methods for detection of nucleic acids
Patent: WO 0190417-A 37 29-NOV-2001;
Marshall, David J. (US) ; Prudent, James R. (US) ; Scherrill,
Christopher B. (US) ; Shapiro, Gideon (US) ; Grenier, Jennifer K.
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                                                                             Marshall,D.J., Prudent,J.R., Scherrill,C.B., Shapiro,G., Grenier,J.K. and Richmond,C.S.
                                                                                                                                                                                                             Sequence 37 from Patent W00190417.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transformation systems for the yeast candida utilis and the expression of heterologous genes therewith Patent: US 5849524-A 23 15-DEC-1998;
                                                                                                                                 artificial
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AR065336
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                                                                                                                                                             synthetic construct.
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   (US) ; Richmond, Craig S. (US)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (bases 1 to 31)
ndo, K., Kajiwara, S. and Misawa, N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 30)
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12 c 5 g
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7 c 9 g
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Pred. No. 16
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                                                                                                                                                                                                      35 bp
Sequence 7 from patent US 5776689
AR016271
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                                                            Protein recruitment system Patent: US 5776689-A 7 07-
                                                                                        1 (bases 1 to 35)
Karin, M., Elledge, S.J. and Aronheim, A.
                                                                                                                                                                                        AR016271.1 GI:3972548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Other publication AU 6805694 950117.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Filipowicz, W. and Connelly, S. CHIMERIC PLANT GENES POSSESSING INDEPENDENT REGULATORY SEQUENCES PATENT: WO 9500652-A 15 05-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           unidentified
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/organism="unknown"
12 c 12 g
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/mod_base=OTHER
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RESULT 85
AR016273/c
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                                                                                                                                                                                  Vector for protein surface presentation Patent: JP 2000189167-A 4 11-JUL-2000; SUNITONO ELECTRIC IND LTD OS ARTIficial Sequence PN JP 2000189167-A/4 PD 11-JUL-2000 PF 28-DEC-1998 JP 1998371747
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Karin,M., Elledge,S.J. and Aronheim,A. Protein recruitment system Patent: US 5776689-A 9 07-JUL-1998;
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                                                                                                                                                                                                                                                                                        Nakajima, H. and Nibara, N.
                                                                                                                                                                                                                                                                                                                   artificial sequences.
                                                                                                                                                                                                                                                                                                                                  synthetic construct. synthetic construct
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JP 2000189167-A/4.
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                                                                                                                                                         HIROTO NAKAJIMA, NAOKI NIBARA
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          /organism="synthetic construct"
/db_xref="taxon:32630"
                                                      Location/Qualifiers
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14 c
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Pred. No. 1
                                                                    /organism='Artificial Sequence'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 15.7%;
Local Similarity 62.2%;
                                                                                               13 GCCGCCGGAGAACATTGAGAT 33
                                                                                                                           72 GCCTCCAGAAAACATTGAGGT 92
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Sequence 9 from patent US 6387619.
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 80 from patent US 6297048. AR171554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chenchik, A., Zhu, Y., Diatchenko, L. and Siebert, P. Methods and compositions for full-length cDNA Cloning using a template switching oligonucleotide patent: US 5962272-A 100 05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 100 from patent US 5962272.
                                                                                                                                                                                                                                                                                              Patent: US 6297048-A 80 02-OCT-2001;
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                   AR209865
                                                                                                                                                                                                                                                                                                           1 (bases 1 to 40) Jolly,D.J., Chang,S.M.W., Lee,W.T.L., Townsend,K. and O'Dea,J. Hepatitis therapeutics
                                                                                                                                                                                                                                                                                                                                                         Unclassified.
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                                                                                                                                                                                                                               12 ACACTATAGGCATGAAGCTGGCATTCCTC 40
                      1 (bases 1 to 40)
Gottschling, D. E. and Singer, M. S.
Yeast telomerase compositions
Patent: US 5698686-A 9 16-DEC-1997;
                                                                                                                                                                                                                                                                               20; Conservative
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Human cystine knot polypeptide
Patent: WO 0153346-A 13 26-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13 from Patent W00153346.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telomerase compositions and methods Patent: US 6387619-A 9 14-MAY-2002;
                                                                                                   Unknown
                                                                                                                           I85801.1 GI:3205519
                                                                                                                                                  Sequence 9 from patent US 5698686.
                                                                                                                                                                185801
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AX201009
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6 GACTGTCGTCCGTCAAAAAATTTCAAGG 34
                                                                                                                                                                                                                                                             Xu,J., Dillon,D.C., Mitcham,J.L., Harlocker,S.L., Jiang,Y., Reed,S.G., Kalos,M.D., Fanger,G.R., Day,C.H., Retter,M.W., Stolk,J.A., Skeiky,Y.A., Wang,A. and Meagher,M.J. Compositions and methods for the therapy and diagnosis of prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AR003527.1 GI:3964786
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                                                                                                                                                                                                      CORIXA CORPORATION (US)
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Fraiser,M.S., Spargo,C.A., Walker,G.Terrance., Van Cleve,M.,
                                                                                                                                                                                                                                                                                                                                             Mammalla; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 45)
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                                                                                                                                                                                                                        Patent: WO 0151633-A 639 19-JUL-2001;
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HSTCELL13/c
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                                                                                                            34 ATGCCTTGGTTTCAACACTATCGGAATGTGACTGTCG 70
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                                                                       39 ATGCCTCCGTGTAAGCCCGAGTAGATTGGGACTATTG
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                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 45)
Fraiser,M.S., Spargo,C.A., Walker,G.Terrance., Van Cleve,M., Wright,D.James. and Little,M.C.
Strand displacement amplification using thermophilic enzymes
Patent: US 5648211-A 4 15-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AGAATGCCTACCGTGCTGCAGTGCGTGAACGTGTCGGTGGTGTCT 45
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Sequence 782 from Patent W00173032.
AX267808
                                                                                                                                                     23;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Unclassified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            I55640.1 GI:2476434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4 from patent US 5648211.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent: WO 0173032-A 782 04-OCT-2001;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                             /organism="unknown"
12 c 10 g
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                                                                                                        Match 15.7%;
Local Similarity 57.8%;
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                                                   35 TGCCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      42 GATGAGATTTCCTTGGCTTCCTCCATATC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27 GGTGACAATGCCTTGGTTTCAACACTATC 55
                3 TGGTTGTGTTAGTTCTCTCTAAAAATGTGGCTGCAGAGCCTGCTG 47
                                                                                          26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 187 from patent US 5833975
AR052850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vie,H., Kabelitz,D. and Bonneville,M.
Surface expression of functional T cell receptor chains formed by interlocus recombination on human T lymphocytes
                                                                                                                                                                                                                                                               Patent: US 5833975-A 187 10-NOV-1998;
                                                                                                                                                                                                                                                                                                                      Paoletti, E., Tartaglia, J. and Cox, W.I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7964454
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H.sapiens rearranged TCR alpha chain variable region (Val3.1Ja42).
                                                                                                                                                                                                                                                                                                  Canarypox virus expressing cytokine and/or tumor-associated antigen
                                                                                                                                                                                                                                                                                                                                                           Unclassified
                                                                                                                                                                                                                                                                                                                                                                                                Unknown.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Direct Submission
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                                                                                          Conservative
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9 c 14 q
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/note="contains part of the V, and J region"
6 c 15 g 12 t
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                                                                                                          Score 14.6; DB 6; Length 47 Pred. No. 1.1e+06;
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             48 ACACTATCGGAATGTGACTGTCGGGCCTC 76
                                                                                                                                                                                                                                                                                                                                                                                                        51 CTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                          38 CTATCGGCAGTCGACAGTCGGCCAGCAAG 10
                                                                                                                                                               Frey,T.K., Dominguez,G. and Wang,C.-Y. DNA encoding infectious rubella virus Patent: US 5439814-A 7 08-AUG-1995;
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Shimizu, A. and Koga, S.
FRUCTOSYL ANINE OXIDASE-PRODUCTIVE SUBSTANTIALLY PURE ORGANISM PAtent: JP 1998201473-A 3 04-AUG-1998;
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20-JAN-1997 JP 1997007101
SHIMIZU AKIRA, KOGA SHINJI
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JP 1998201473-A/3
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Location/Qualifiers
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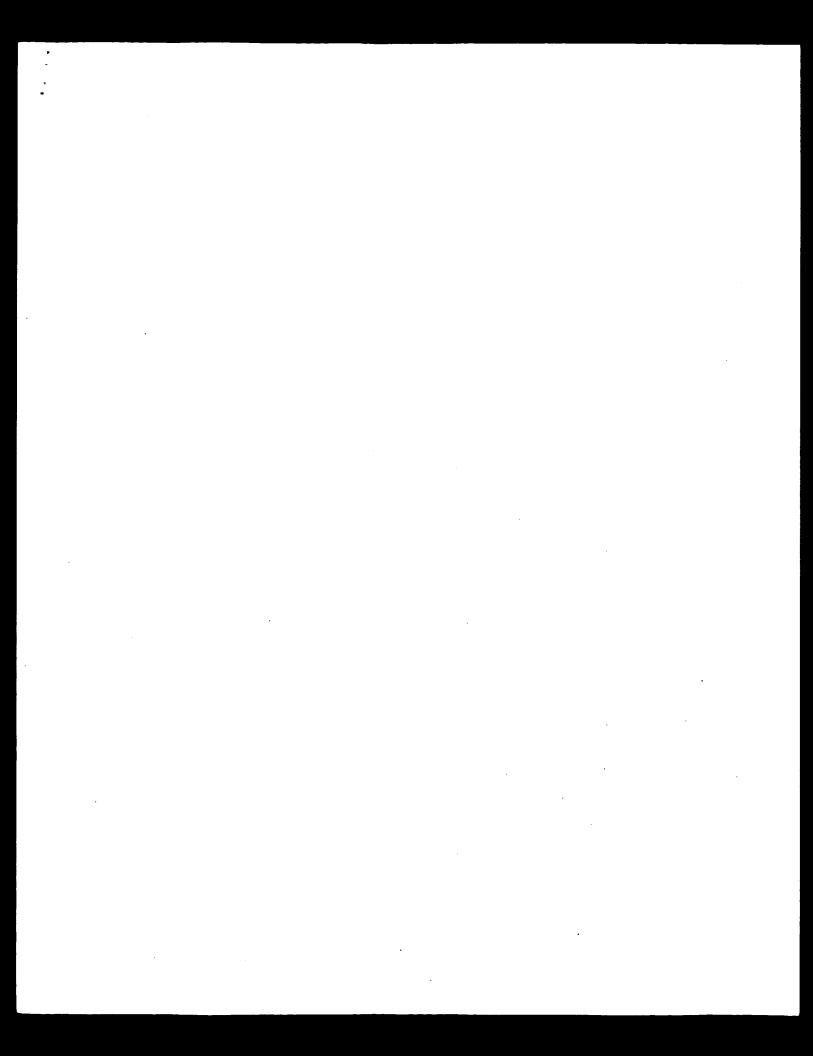
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                                        20 ACACTATAGCATGGAAGCTATCGGACCTC 48
                                                                   48 ACACTATCGGAATGTGACTGTCGGGCCTC 76
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Frey, F.K., Dominguez, G. and Wang, C.-Y.
DNA encoding infectious Rubella virus
Patent: US 5663065-A 4 02-SEP-1997;
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163473
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Sequence 45, Appl
Sequence 5613, Ap
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Sequence 40, Appl
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SOFTWARE: Microsoft Office 97
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LENGTH: 31
PRIOR FILING DATE: 2000-11-14
                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence:Synthetic ; OTHER INFORMATION: Oligonucleotide US-09-993-170-22
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US-09-993-170-22
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APPLICANT: Bryan, Gregory
TITLE OF INVENTION: Modification of a Plant Disease Resistance Gene
TITLE OF INVENTION: Specificity and Method for Engineering Altered
TITLE OF INVENTION: Specificity
FILE REFERENCE: BB135 US NA
                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/993,170 CURRENT FILING DATE: 2001-11-26
                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           37 CCTTGGTTTCAACACTATCGGAATGTGAC 65
                                                                                                         21; Conservative
                                                  CCTGGGATCCAACACTAACGTTATTTAAC 30
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US-0016-17A-440
US-09-998-30A-440
US-09-998-757A-440
US-09-978-757A-440
US-09-978-187B-440
US-09-978-187B-440
US-09-978-643A-440
US-09-978-643A-440
US-09-978-643A-440
US-09-978-643A-440
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0 US-09-263-959-143
US-09-992-598-385
US-09-989-735-385
US-09-989-735-385
US-09-989-730-385
US-09-991-181-385
US-09-991-181-385
US-09-991-181-385
US-09-991-653-385
US-09-991-67-385
US-09-991-67-385
US-09-991-67-385
US-09-991-67-385
US-09-991-67-385
US-09-991-68-385
US-09-991-18-385
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                                                                                                         Mismatches
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                                                                                                                                   DB 10;
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                                                                                                                                   Length 31;
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Sequence 440, App
Sequence 10, Appl
Sequence 113, Appl
Sequence 313, App
Sequence 385, App
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Sequence 440, App
Sequence 440, App
Sequence 440, App
Sequence 14, Appl
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Sequence
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                                                                                                      Gaps
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                                                                                                        0;
                                          CURRENT APPLICATION NUMBER: US/09/801,371A
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: PCT WO 00/14255
PRIOR FILING DATE: 1999-09-06
PRIOR FILING DATE: 1999-09-06
NUMBER OF SEO ID NOS: 12
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 43
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-801-371A-2
                                                                                                                                                                                                                                                                            ; Sequence 2, Application US/09801371A
; Patent No. US20020155569A1
                                                                                                                                                                                                                                                                                                         US-09-801-371A-2
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                                                                                                                                                                                                       APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                            TITLE OF INVENTION: REGULATION OF GENE EXPRESSION THROUGH TITLE OF INVENTION: MANIPULATION OF MRNA SPLICING AND ITS USES FILE REFERENCE: A34084-PCT-USA-A 066031.0147
                                                                                                                                                                                                                               APPLICANT: Kaempfer, Raymond
APPLICANT: Osman, Farhat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences NUMBER OF SEQUENCES: 5255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: HP Vect
OPERATING SYSTEM:
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CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
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Local Similarity 72.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50 base pairs
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Pred. No. 1.1e+04;
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Matches

RESULT 2

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; ORGANISM: Homo sapien US-09-801-371A-8
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                                                                                                                                                                                                                                       SEQ ID NO 8
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                                                                   Matches
                                                                                                         Query Match
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                                                                                         Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Jarrous, Nayef
APPLICANT: Ben-Ascoli, Yitzhak
TITLE OF INVENTION: REGULATION OF GENE EXPRESSION THROUGH
TITLE OF INVENTION: MANIPULATION OF MRNA SPLICING AND ITS USES
FILE REFERENCE: A34084-PCT-USA-A 066031.0147
CURRENT APPLICATION NUMBER: US/09/801,371A
CURRENT FILING DATE: 2001-03-07
                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Jarrous, Nayef
APPLICANT: Ben-Asouli, Yitzhak
TITLE OF INVENTION: REGULATION OF GENE EXPRESSION THROUGH
TITLE OF INVENTION: MANIPULATION OF MRNA SPLICING AND ITS USES
FILE REFERENCE: A34084-PCT-USA-A 066031,0147
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PRIOR APPLICATION NUMBER: PCT WO 00/14255
PRIOR FILING DATE: 1999-09-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kaempfer, Raymond
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                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Homo sapien
                                                                                                                                                                                                TYPE: RNA
                                                                                                                                                                                                                     CENGTH:
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                     68 TCGGGCCTCCAGAAAACATTGAGG 91
                                                                   16; Conservative
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  Osman, Farhat
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79.2%;
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66.7%;

 Mismatches

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                                                                                    Score 16; DB 9;
Pred. No. 1.3e+04;
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Pred. No. 1.
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                                                                                                       DB 9; Length 50;
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US-10-043-573-45/c
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Publication No. US20020182724A1
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 78554
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                                                                                                                                                                        Matches
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APPLICANT: Friedrich, Glenn A.
APPLICANT: Liliaberg, Stan
APPLICANT: Liliaberg, Stan
TITLE OF INVENTION: VECTORS FOR GENE MUTAGENESIS AND GENE
TITLE OF INVENTION: DISCOVERY
FILE REFERENCE: 7705-0006-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Mittman, Michael TITLE OF INVENTION: Human MFILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/081,727
PRIOR FILING DATE: 1998-04-14
PRIOR APPLICATION NUMBER: US 60/109,302
PRIOR FILING DATE: 1998-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/158,735
CURRENT FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 09/276,533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/079,729 PRIOR FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1999-03-25
                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                               ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                         TYPE: DNA
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                                                                                                                            67 GTCGGGCCTCCAGAAAACATTGAGGTG 93
                                                                                                                                                                      20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  66 TGTCGGGCCTCCAGAAAAC 84
                                                                               33 GGCGGTCCTGCAGGGAACAGTGGGGTG 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 UGGGGCCUCCAGAACUCACUGGGG 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 TGTCGGTCCTCCAGAAGAC 20
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                                                                                                                                                                                           Similarity
                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                           ID NOS:
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                                                                                                                                                                                           17.0%; Score 15.8; DB 9; Length 39; 74.1%; Pred. No. 1.4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17.0%; Score 15.8; DB 9; Length 25; 89.5%; Pred. No. 1.2e+04;
                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                      0,
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SEQUENCE DESCRIPTION: SEQ ID NO: 45:
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                                                                                                                                                                                             Sequence 1028, Application US/10215112 Publication No. US20030082596A1 GENERAL INFORMATION:
SEQ ID NO 1028
                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                        TITLE OF INVENTION: Method of Genetic Analysis of Probes:
TITLE OF INVENTION: Test3
FILE REFERENCE: 3119
                CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                             APPLICANT: Michael Mittmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 45:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 018547-030100US TELECOMMUNICATION INFORMATION: TELEPHONE: 415 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/813,507

FILING DATE: 07-MAR-1997

APPLICATION NUMBER: US 60/032,069

FILING DATE: 02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sapolsky, Ronald J. TITLE OF INVENTION: Brassica Polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lemieux, Bertrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                             43 TTTCAACACTATCGGAATGTGACTGTC 69
                                                                                                                                                                                                                                                                                                                                                     32 TTGCGACTCTGTGGGACTGTAACTGTC 6
                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: replace(21, "")
OTHER INFORMATION: /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/10/043,573 FILING DATE: 09-Jan-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Liebeschuetz, Joe
REGISTRATION NUMBER: 37,505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 415 576-0200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NDDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Landry, Benoit S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -3834
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74.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.8; DB 9;
Pred. No. 1.5e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 9; Length 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                           7; Indels
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; ORGANISM: Homo sapiens US-09-801-274-694
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                                                                                                                                          SOFTWARE: Fas
SEQ ID NO 694
LENGTH: 31
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        Query Match
Best Local Similarity
Matches 17; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 694, Application US/09801274
Patent No. US20020032319A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 5613
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                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Cargill, Michele APPLICANT: Ireland, James : APPLICANT: Lander, Eric S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Method of Genetic Analysis of Probes:
TITLE OF INVENTION: Test3
FILE REFERENCE: 3119
CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
NUMBER OF SEQ ID NOS: 14936
SOFTMARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/206,129 PRIOR FILING DATE: 2000-05-22
                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/187,510 PRIOR FILING DATE: 2000-03-07
                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
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TYPE: DN
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                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 25
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30 GACAATGCCTTGGTTTCAACAC 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 GTCGATTCCTGGGTTTCAACAC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                    FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                   Ireland, James S.
          Conservative
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                           16.3%;
77.3%;
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81.8%;
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    Mismatches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                         Score 15.2; DB 10;
Pred. No. 2.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.6; DB 9; Length 25; Pred. No. 1.5e+04;
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Pred. No. 1.
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Indels
                                              Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
          0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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; OTHER INFORMATION: Primer Used in Examples 5C and 5D US-09-231-235-40
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US-09-231-235-40
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                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 40, Application US/09231235
Patent No. US20020048805A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 40
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             Matches
                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/231,235
CURRENT FILING DATE: 1999-01-15
NUMBER OF SEQ ID NOS: 63
                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: FELINE IMMUNODEFICIENCY VIRUS GENE THERAPY VECTORS FILE REFERENCE: 930049.467
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Sheridan, Philip Lee
APPLICANT: Hardy, Steven
APPLICANT: Dubensky, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Sauter, Sybille L. APPLICANT: Hsu, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/872,696A CURRENT FILING DATE: 1999-01-18 NUMBER OF SEQ ID NOS: 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Dubensky, Thomas
APPLICANT: Yee, Jiing-Kuan
TITLE OF INVENTION: FELINE IMMUNODEFICIENCY VIRUS GENE THERAPY VECTORS
FILE REFERENCE: 930049.467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Johnston, Julie C.
APPLICANT: Sauter, Sybille L
APPLICANT: Hsu, David
APPLICANT: Sheridan, Philip
APPLICANT: Hardy, Steven
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Yee, Jiing-Kuan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Johnston, Julie C.
                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Primer OTHER INFORMATION: Primer Used in Examples 5C and 5D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                               LENGTH: 46
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             20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 CTCCATTCTGCCTGGGTGACAATGCCTTGG 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 CCATTCTGCCTGGGTGACAATG 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13 CCCCATCCTACCTTGTYGACTGTCCCTCGG 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 CCGCTCWGCCTGGATGCCAATG 1
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Dubensky, Thomas
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           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
    16.3%; Score 15.2; DB 10; Length 46; 66.7%; Pred. No. 2.6e+04; ative 1; Mismatches 9; Indels
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    Gaps
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    0;
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                                                                 ; OTHER INFORMATION: Annealing oligonucleotide US-09-935-280-15
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US-09-935-280-15
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                                                                                                                                                                         SOFTWARE: PatentIn version 3.0 SEQ ID NO 15 LENGTH: 15
Best Local Similarity
                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15, Application US/09935280 Publication No. US20030022179A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 40
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                                                                                                                                                                                                                                                                                               FILE REFERENCE: INVITI300-1
CURRENT APPLICATION NUMBER: US/09/935,280
CURRENT FILING DATE: 2001-08-21
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: MADDEN, Knut
APPLICANT: BENNETT, ROB
TITLE OF INVENTION: METHODS AND REAGENTS FOR MOLECULAR CLONING
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: INVITROGEN CORPORATION APPLICANT: CHESTNUT, John APPLICANT: SHUMAN, Stewart APPLICANT: HEYMAN, John
                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/226,563 PRIOR FILING DATE: 2000-08-21
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                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 18
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CURRENT FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: 09/231,235
DECOS STITES AND ASSESSED 117
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TITLE OF INVENTION: FELINE IMMUNODEFICIENCY VIRUS GENE THERAPY VECTORS
FILE REFERENCE: 930049.467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Johnston, Julie C. APPLICANT: Sauter, Sybille L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1999-01-15
NUMBER OF SEQ ID NOS: 63
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                                                                                                                                                  TYPE: DNA
                                                                                                             FEATURE:
                                                                                                                               ORGANISM: Artificial sequence
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16.1%; Score 15; DB 9; I 100.0%; Pred. No. 2.2e+04;
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                    Length 15;
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US-09-792-875-1
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US-09-792-875-1
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NUMBER OF SEQ ID NOS: 21
SOFTWARE: Patentin version 3.0
SEQ ID NO 1
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Patent No. US20010044137A1
GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                     Query Match
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APPLICANT: HEYMAN, John
APPLICANT: SZALAI, Gabor
TITLE OF INVENTION: TOPOISOMERASE LINKER-MEDIATED AMPLIFICATION METHODS
FILE REFERENCE: INVIT1270-1
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PRIOR FILING DATE: 1998-04-03
NUMBER OF SEQ ID NOS: 6
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PRIOR FILING DATE: 1999-04-02
PRIOR APPLICATION NUMBER: US 60/096,981
PRIOR FILING DATE: 1998-08-18
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CURRENT FILING DATE: 2001-11-14
                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/792,875
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TITLE OF INVENTION: LIBRARIES OF EXPRESSIBLE GENE SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: INVITROGEN CORPORATION APPLICANT: FERNANDEZ, Joseph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: HEYMAN, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial sequence
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                                                                                                                                                                   OTHER INFORMATION: oligonucleotide Topo5
                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                       ORGANISM: ARTIFICIAL
                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                     ENGTH: 16
                                                         Local Similarity es 15; Conserv
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                  46 CAACACTATCGGAAT 60
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                                                  16.1
100.08; F1
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100.0%; Pred. No. 2.2e+04;
                                                                              16.1%; score 15; DB 10; Length 16; 100.0%; Pred. No. 2.2e+04;
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                                                           Mismatches
                                                           2.2e+04;
hes 0;
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                                                           0
                GEMERAL INFORMATION:
APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
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APPLICANY: Mittman, Michael TITLE OF INVENTION: Human Microarray FILE REFERENCE: 3118.1
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                                                                                                                                                                ; Sequence 86605, Application US/10098263B; Publication No. US20030104410A1
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PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
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PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
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PRIOR FILING DATE: 2001-03-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapien
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                                                                                                                                                                                                                                                                                                                                                                                   0;
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; TYPE: DNA; ORGANISM: Homo sapiens US-09-801-274-1732
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-801-274-1732/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-098-263B-116206
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 116206
                                                                                                                                                                 NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1732, Application US/09801274 Patent No. US20020032319A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 86605
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                                            Query Match
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      Matches
                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR FILING DATE: 2000-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR APPLICATION NUMBER: US 60/187,510
                                                                                                                                                                                                                                                                                                        APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cargill, Michele APPLICANT: Ireland, James S. APPLICANT: Lander, Eric S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILLING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILLING DATE: 2001-03-16
PRIOR FILLING DATE: 2001-03-16
                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/206,129 PRIOR FILING DATE: 2000-05-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapien
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ORGANISM: Homo sapien
                                                                                                                                               LENGTH:
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                      Match 16.1%;
Local Similarity 72.0%;
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Local Similarity 78.3%;
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  18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24 CGGACCTCCGGAAAACACGGACG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      69 CGGGCCTCCAGAAAACATTGAGG 91
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18; Conservative
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Similarity 78.3%;
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    Mismatches

                   Score 15; DB 10;
Pred. No. 2.7e+04;
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Pred. No. 2.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15; DB 9;
Pred. No. 2.5e+04;
                                          DB 10; Length 31;
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  Indels
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                   RESULT 24
US-10-112-802-60/c
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; Sequence 60, Application US/10112802
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                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/900,379
FILING DATE: 06-Jul-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                      SEQUENCE DESCRIPTION: SEQ ID NO: 38:
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (708) 267-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 01-FEB-1995
APPLICATION NUMBER: US 08/283,302
FILING DATE: 29-UUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jiao, Jin-An
TITLE OF INVENTION: MHC COMPLEXES AND USES THEREOF
NUMBER OF SEQUENCES: 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Wong, Hing C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                         11 CACTCCATTCTGCCTGGGTGACAATGCCTTG 41
                                                                                                                                                                                21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28 GTGACAATGCCTTGGTTTCAACACT 52
                                                                                                   1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US95/09816 FILING DATE: 31-JUL-1995 APPLICATION NUMBER: US 08/382,454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Deerfield
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/776,084 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Dade International, Inc
STREET: 1717 Deerfield Road
                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 32,369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Pearson, Louise S.
                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                               relefax: (708)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chavaillaz, Pierre-Andre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Widanz, Jon A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rhode, Peter R.
                                                                                                                                                                                                  16.1%; Score 15; DB 9; Length 42; 67.7%; Pred. No. 3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                               267-5376
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US-10-112-802-61
US-10-112-802-61, Application US/10112802
; Sequence 61, Application US/10112802
; Publication No. US20030086942A1
                                                                                                      US-09-848-164-38
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                        Sequence 38, Application US/09848164 Patent No. US20020034513A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/280,214
PRIOR FILING DATE: 2001-03-30
NUMBER OF SEQ ID NOS: 75
SOFTMARE: Patentin Ver. 2.1
SEQ ID NO 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/280,214
PRIOR FILING DATE: 2001-03-30
NUMBER OF SEQ ID NOS: 75
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Publication No. US20030086942A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Petersen, Jeannine M. APPLICANT: Her, Lu-Shiun APPLICANT: Dahlberg, James E. APPLICANT: Glodowski, Doreen R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/112,802
CURRENT FILING DATE: 2002-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Inhibition of Nucleocytoplasmic Transport by Vesicular TITLE OF INVENTION: Stomatitis Virus M Protein-Like Polypeptides FILE REFERENCE: 960296.97770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 960296.97770
CURRENT APPLICATION NUMBER: US/10/112,802
CURRENT FILING DATE: 2002-03-29
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APPLICANT: Glodowski, Doreen R.
TITLE OF INVENTION: Stomatitis Virus M Protein-Like Polypeptides
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APPLICANT: Her, Lu-Shiun
APPLICANT: Dahlberg, James E.
APPLICANT: Glodowski, Doreen R.
                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence:oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                                                                                                                                                                                                                                                        Match 16.1%; Score 15; DB 9; Local Similarity 61.5%; Pred. No. 3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
APPLICANT: Rhode, Peter R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           42 GGGAGGGGTCTTCCCCATTCTGGCTGGCAAATAAGCCCT 4
                                                                                                                                                                                                                                                                                                     24; Conservative
                                                                                                                                                                                                                                               1 GAGCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                 1 GGGAGGGGTCTTCCCCATTCTGGCTGGCAAATAAGCCCCT 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16.1%; Score 15; DB 9; Length 42; 61.5%; Pred. No. 3e+04;
                                                                                                                                                                                                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                DB 9; Length 42;
                                                                                                                                                                                                                                                                                                     15; Indels
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; Sequence 7289, Application US/10156306
; Publication No. US20030119017A1
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                                                                                                                                                                                                                                                                                                                                                                    ; GENERAL INFORMATION:
                                                                                                             TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Rel TITLE OF INVENTION: Levels of IKK-Gamma and PKR FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTMARE: Patentin version 3.0
SEQ ID NO 7289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                  APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: McSwiggen, James
OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic Acid
                                         ORGANISM: Artificial Sequence
                                                                      TYPE: RNA
                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 38:
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FILING DATE: <UNKNOWN>
APPLICATION NUMBER: US 08/382,454
FILING DATE: 01-FEB-1995
APPLICATION NUMBER: US 08/283,302
FILING DATE: 29-UL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE DESCRIPTION: SEQ ID NO: 38:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wong, Hing
TITLE OF INVENTION: MHC COMPLEXES AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11 CACTCCATTCTGCCTGGGTGACAATGCCTTG 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tch 16.1%;
al Similarity 67.7%;
21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/848,164 FILING DATE: 03-May-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Deerfield
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (708) 267-5300
TELEFAX: (708) 267-5376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Pearson, Louise S. REGISTRATION NUMBER: 32,369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Dade International, Inc. STREET: 1717 Deerfield Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: STR-4665-CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jiao, Jin-An
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            124
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US-10-156-306-7289

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; OTHER INFORMATION: Description of Artificial Sequence:PRIMER US-09-245-764-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1 SEQ ID NO 23348
      Best Local Similarity 73. Matches 19; Conservative
                                        Query Match
                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/09245764 Patent No. US20020107359A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                      FILE REFERENCE: 4102-4
CURRENT APPLICATION NUMBER: US/09/245,764
CURRENT FILING DATE: 1999-02-05
EARLIER APPLICATION NUMBER: 60/099,994
EARLIER FILING DATE: 1998-09-11
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Epa, Vidana
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS
TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                              EARLIER APPLICATION NUMBER: 60/073,972
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: POWell, Maree S.
APPLICANT: McKenzie, Ian F.C.
APPLICANT: Maxwell, Kelly F.
APPLICANT: Garrett, Thomas P.J.
                                                                                                                 TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                           ENGTH: 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18 TTCTGCCTGGGTGACAATGCCTT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 TGACCGTCGGGTCTCCAG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hogarth, P. Mark
                      15.9%;
73.1%;
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      0; Mismatches
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Pred. No.
                      Score 14.8; DB 10; Length 30; Pred. No. 3.2e+04;
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Pred. No. 3e
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      Indels
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Gaps
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US-09-780-533A-3768
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-912-263-485/c
                                                                                                                                                                                                                                                                                                                                                                      ; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 3768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; SEQ ID NO 485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; GENERAL INFORMATION:
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                                                                                    Matches
                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: Method, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780.533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR PILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric. S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2017-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Blatt, Larry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/912,263
CURRENT FILING DATE: 2001-07-24
PRIOR APPLICATION NUMBER: US 60/220,315
PRIOR FILING DATE: 2000-07-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Cargill, Michele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE:
                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic Acid NAME/KP: misc_feature LOCATION: (31) (31) OTHER INFORMATION: n stands for inosine
                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                              TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                                           Local
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                                          51 CTATCGGAATGTGACTGTCGGGCCTCCAGAAAACA 85
                                                                                    18;
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4 CUAUCCUGAUGAGGCCGUUAGGCCGAANCAAAACA 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28 CTGGCTGCAATCSATTCTGC 9
                                                                                                                                                                                                                                                                                                                                                   38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 CTGGGAGCACTCCATTCTGC 23
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                                                                                                         h 15.9%;
Similarity 51.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chowrira, Bharat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McSwiggen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/09912263
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                                                                                    Conservative
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                                                                                    4; Mismatches
                                                                                                         Score 14.8; DB 9;
Pred. No. 3.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                              Length 38;
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                                                                                    Indels
                                                                                    0;
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                                                                                 Gaps
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RESULT 33
US-10-060-275-7
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US-10-043-573-147
                                                                                                                                                                                                                     Sequence 7, Application US/10060275 Publication No. US20030073828A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
APPLICANT: DAI, ZIYU
APPLICANT: SHI, LIFANG
APPLICANT: HOOKER, BRIAN S.
TITLE OF INVENTION: EPIMERASE GENE AND USE THEREOF
FILE REFERENCE: 059440-0143
CURRENT APPLICATION NUMBER: US/10/060,275
CURRENT FILING DATE: 2002-06-11
PRIOR APPLICATION NUMBER: 60/265,311
PRIOR FILING DATE: 2001-02-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 147:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION
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nes 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/813,507
FILING DATE: 07-MAR-1997
APPLICATION NUMBER: US 60/032,069
FILING DATE: 02-DEC-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 415 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sapolsky, Ronald J. TITLE OF INVENTION: Brassica Polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lemieux, Bertrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                         28 GTGACAATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                               6 GTGAAAAGGCATTGTKCTCCAAAATCTCGCTCTGCG 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 41 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/043,573 FILING DATE: 09-Jan-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Liebeschuetz, Joe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 415 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 37,505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Landry, Benoit S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15.9%; Score 14.8; DB 9; Length 41; 61.1%; Pred. No. 3.5e+04;
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US-09-904-380-12/c
; Sequence 12, Application US/09904380
; Patent No. US20020022229A1
                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 4, Application US/10076631; Publication No. US20030028001A1; GENERAL INFORMATION:
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                                                                                      us-10-076-631-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-076-631-4
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                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 4
LENGTH: 46
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  Matches
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Best Local Similarity 64.7%;
Matches 22; Conservative
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                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Use of supersecretable peptides in processes for their TITLE OF INVENTION: production and for parallel improvement of the exported TITLE OF INVENTION: forms of one or more other polypeptides of interest FILE REFERENCE: DEAV2001/0007
                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/076,631
CURRENT FILING DATE: 2002-02-19
NUMBER OF SEQ ID NOS: 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Aventis Pharma Deutschland GmbH
                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                      OTHER INFORMATION: Description of Artificial Sequence:hirfl
                                                                                                                                                                      TYPE: DNA
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                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 CTGCCTGGGTGACAATGCCTTGGTTT 45
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Similarity 73.1%;
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  Conservative
                  15.9%; Score 14.8; DB 9; 64.7%; Pred. No. 3.7e+04;
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Pred. No. 3.6e+04;
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Pred. No. 3.5e+04;
  Mismatches
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                                          DB 9; Length 46;
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34 ATGCCTTGGTTTCAACACTATCGGAATGTGACTG 67

10 ATCCTTTGGATAAAAGACTTACGTATACTGACTG 43

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Publication No. US20030082596A1
GENERAL IMPORMATION:
APPLICANT: Michael Mittmann
GENERAL INFORMATION:
APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1 SEQ ID NO 40831
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                                                                                                                                   Sequence 99102, Application US/10098263B Publication No. US20030104410A1
                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Method of Genetic Analysis of Probes: TITLE OF INVENTION: Test3
FILE REFERENCE: 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Mittman, Michael TITLE OF INVENTION: Human Microarray FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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Local Similarity 81.0%;
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Local Similarity 81.0%;
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                                                                                                                                                                                                                                                                                                                 24 CTGGGTGACAATGCCTTGGTT 44
                                                                                                                                                                                                                                                                      17; Conservative
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                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                               Score 14.6; DB 9; Pred. No. 3.6e+04;
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Pred. No. 3.
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; GENERAL ; APPLIC!

APPLICANT: KONDO, KEIJI

; Sequence 30, Application US/09908855; Patent No. US20020115220A1; GENERAL INFORMATION:

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RESULT 40
US-09-908-855-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/205,712
PRIOR FILING DATE: 2000-05-19
PRIOR APPLICATION NUMBER: US 60/240,398
PRIOR FILING DATE: 2000-10-14
PRIOR APPLICATION NUMBER: US 60/282,831
PRIOR FILING DATE: 2001-04-10
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CURRENT FILING DATE: 2001-05-18
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APPLICANT: Prudent, James R.
APPLICANT: Sherrill, Christo
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TITLE OF INVENTION: MATERIALS AND METHODS FOR DETECTION OF NUCLEIC ACIDS
FILE REFERENCE: 031403-9006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                          OTHER INFORMATION: n represents deoxy-iso-cytosine (d-isoC)
NAME/KEY: modified_base
LOCATION: (16)..(16)
OTHER INFORMATION: n represents deoxy-iso-guanine (d-isoG)
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                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: modified_base LOCATION: (15)..(15)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: oligonucleotide FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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                                                                                                                                                                  15.7%; Score 14.6; DB 10 Local Similarity 73.9%; Pred. No. 3.9e+04; es 17; Conservative 0. 16.2.1
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                                                                                     24 CAGGGATCNNTCCTGTCTGCCTG 2
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                                                                                                                             4 CTGGGAGCACTCCATTCTGCCTG 26
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Sherrill, Christopher B.
Shapiro, Gideon
Grenier, Jennifer K.
Richmond, Craig S.
Jurczyk, Simona
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17; Conservative
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 Mismatches

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JU-181-836-13

Sequence 13, Application US/10181836
Publication NO. US20030059877A1
GENERAL INFORMATION:
APPLICANY: AKZO NO. USC.
FILE REFT.
                                                                                                                                                                                                                                                    RESULT 42
US-09-912-679-80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-181-836-13
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                                                                                                                                                                                 Sequence 80, Application US/09912679 Patent No. US20020141974A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/EP01/00571
PRIOR FILING DATE: 2001-01-17
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PATENTIN VETSION 3.1
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 30
LENGTH: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Akzo No. US20030059877A1e1
TITLE OF INVENTION: NOVEL CYSTINE KNOT POLYPEPTIDE
FILE REFERENCE: 2000.527 us
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/181,836 CURRENT FILING DATE: 2002-07-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/908,855
CURRENT FILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: 09/242,690
PRIOR FILING DATE: 1999-02-23
PRIOR APPLICATION NUMBER: JP 8/241062
PRIOR FILING DATE: 1996-08-23
RIOR FILING DATE: 1996-08-23
NUMBER OF SEQ ID NOS: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MIURA, YUTAKA
TITLE OF INVENTION: YEAST VECTOR AND METHOD OF PRODUCING PROTEINS USING THE
TITLE OF INVENTION: SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 049441/0118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ocal Similarity 81.0
                            TITLE OF INVENTION: HEPATITIS THERAPEUTICS NUMBER OF SEQUENCES: 84
    CORRESPONDENCE ADDRESS:
                                                                                                                                                         APPLICANT: Jolly, Douglas J
                                                                                                                                                                                                                                                                                                                                        12 ACACTATAGGCATGAAGCTGGCATTCCTC 40
                                                                                                                                                                                                                                                                                                                                                                                   48 ACACTATCGGAATGTGACTGTCGGGCCTC 76.
                                                                                                                                                                                                                                                                                                                                                                                                                                 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     40
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                                                                   O'Dea, Joanne
                                                                                                              Chang, Stephen M. Lee, William T.L.
                                                                                           Townsend, Kay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15.7%; Score 14.6; DB 10; Length 38; 81.0%; Pred. No. 4.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                            15.7%; Score 14.6; DB 9;
69.0%; Pred. No. 4.2e+04;
ative 0; Mismatches 9
                                                                                                                                      Stephen M.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 9; Length 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                           ; OTHER INFORMATION: oligo 55 US-10-193-960-55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : Sequence 55, Application US/10193960 : Publication No. US20030104477A1 : GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-193-960-55/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-912-679-80
                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 55
  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: WO PCT/US98/06704 PRIOR FILING DATE: 1998-04-03
                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 08/832,985
PRIOR FILING DATE: 1997-04-04
PRIOR APPLICATION NUMBER: US 08/835,159
PRIOR FILING DATE: 1997-04-04
                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/193,960 CURRENT FILING DATE: 2002-07-12 PRIOR APPLICATION NUMBER: US/09/410,903 PRIOR FILING DATE: 1999-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Biosite Diagnostics Inc.
TITLE OF INVENTION: Chimeric Polyclonal Antibodies
FILE REFERENCE: 014907-002700US
                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 100
                                                                                        FEATURE:
                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 3723836
INFORMATION FOR SEQ ID NO: 80:
                                                                                                                                                       LENGTH: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 15.7%;
Local Similarity 81.0%;
les 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/912,679
FILING DATE: 07-Jun-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 98104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 80:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13 GCCGCCGGAGAACATTGAGAT 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        72 GCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 930049.407C5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: McMasters, David D. REGISTRATION NUMBER: 33,963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Seed and Berry
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15.7%; Score 14.6;
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Pred. No. 4.2e+04;
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  DB 9;
Length 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 40;
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Gaps

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                                                                                                                                                                                                                US-09-895-793-782
                                                                                                                                                                                                                                 RESULT 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-012-896-782
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                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 782
LENGTH: 45
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                                                                                                                                                                            Sequence 782, Appublication No.
                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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APPLICANT: Dill
APPLICANT: Mitc
APPLICANT: Harl
                                                                                                                                                                                                                                                                                                                                                                           Best
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                                                   APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Jiang, Yuqiu
APPLICANT: Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 1011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Meagher, Madeleine Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C27
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 3.0
                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                   31 ACAATGCCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCT 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     39 CCTGGGTTATCACAATATCGGCCATGGCTGGTTGGGC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           37 CCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGC 73
                                                                                                                                                                                                                                                                                1 AGAATGCCTACCGTGCTGCAGTGCGTGAACGTGTCGGTGTCT 45
                                                                                                                                                                                                                                                                                                                                                        26;
                                                                                                                                                                                                                                                                                                                                                                        Similarity
Day, Craig H.
                  Retter, Marc W. Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wantanabe,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Foy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        McNeill, Patricia D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hural, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Retter, Marc W. Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Harlocker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dillon, Davin C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fanger, Gary R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Henderson, Robert A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wang, Aijun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Carter, Darrick
                                                                                                                                                                                         Application US/09895793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10012896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                            US20020192763A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jiangchun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Samuel X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Teresa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craig H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          on, Raymond L.
de Bassols, Carlota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           William T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yasir A.W.
                                                                                                                                                                                                                                                                                                                                                                      15.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yoshihiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Susan L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 4.3e+04;
                                                                                                                                                                                                                                                                                                                                                                    Score 14.6; DB 9; Length 45; Pred. No. 4.4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                       19; Indels
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                                SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 782 LENGTH: 45
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                                                                                   CURRENT APPLICATION NUMBER: US/09/895,814
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 990
                                                                                                                                      FILE REFERENCE: 210121.427C26
                                                                                                                                                     APPLICANT: Fanger, Gary R.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
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APPLICANT: Di
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CURRENT FILING DATE: 2001-06-29
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ORGANISM: Homo sapiens
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Kalos, Michael D.
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Hepler, William T.
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                                                                                                                                                                                                                                              Houghton, Raymond L.
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Hepler, William T.
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57.8%;
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US-09-895-814-782

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ang, Augurant: mang, Aljun

APPLICANT: Skeiky, Yasir A.W.

APPLICANT: Hepler, William

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23

CURRENT APPLICATION NUMBER: US/09/759,143

CURRENT FILING DATE: 2001-01-12

NUMBER OF SEQ ID NOS: 934

SEG ID NO 782

LENGTH: 45

TYPE: 7
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US-09-780-669-782
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US-09-759-143-782
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                                                                                                                                                                                                                                                                           INFORMATION:
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  Carter, Darrick
                                                              Stolk,
                                                                                                                                                                                          Harlocker, Susan L.
                                                                                Fanger, Gary R. Retter, Marc W.
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                      Vedvick, Thomas S.
                                          Day, Craig H.
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                                                              John A.
                                                                                                                                                                    Yugu
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                                                                                                                                                  Robert A.
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; Patent No. US20020098546A1
                                                                                                                                                                                                                                                                                           US-09-918-063-53/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens US-09-822-827-782
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; SEQ ID NO 782
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; APPLICANY: Xu, Jiangchun
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; FILE REFERENCE: 210121.534C1
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-780-669-782
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; Patent No. US20020081680A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity 57.8%;
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                                                                                                                       APPLICANT: Jensen, Wayne A.
APPLICANT: Chandrashekar, Ramaswamy
TITLE OF INVENTION: CANINE TAG1 PROTEINS, NUCLEIC ACID MOLECULES, AND USES THEREOF
FILE REFERENCE: DI-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
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CURRENT APPLICATION NUMBER: US/09/780,669

CURRENT FILING DATE: 2001-02-09

NUMBER OF SEQ ID NOS: 943
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APPLICANT:
               PRIOR APPLICATION NUMBER: 60/224,655
PRIOR FILING DATE: 2000-08-11
NUMBER OF SEQ ID NOS: 85
                                                                              CURRENT APPLICATION NUMBER: US/09/918,063
CURRENT FILING DATE: 2001-07-30
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APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C24
SOFTWARE: PatentIn version 3.1
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Local Similarity 57.8%;
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Hepler, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hural, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wang, Aijun
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.6; DB 10; Length 45; Pred. No. 4.4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.6; DB 10; Length 45; Pred. No. 4.4e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Canis familiaris US-09-918-063-55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-918-063-55
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                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19, Application US/09996073 Publication No. US20030003565A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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LENGTH: 45
                                                                                                                                                                   SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                  APPLICANT: DUBENSKY JR., Thomas W.
APPLICANT: GASMI, Mehdi
APPLICANT: SAUTER, Sybille
APPLICANT: SAUTER, Sybille
TITLE OF INVENTION: FUNCTIONAL LENTIVIRAL VECTOR FROM
TITLE OF INVENTION: AN MLV-BASED BACKBONE
FILE REFERENCE: 2302-1642 / 1642-002
CURRENT APPLICATION NUMBER: US/09/996,073
CURRENT FILING DATE: 2001-11-27
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Weber, Eric R.
APPLICANT: Jensen, Wayne A.
APPLICANT: Chandrashekar, Ramaswamy
TITLE OF INVENTION: CANINE TAGI PROTEINS, NUCLEIC ACID MOLECULES, AND USES THEREOF
FILE REFERENCE: DI-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/224,655
PRIOR FILING DATE: 2000-08-11
NUMBER OF SEQ ID NOS: 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/918,063
CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1
                                                                                                                       LENGTH: 46
TYPE: DNA
                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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LOCATION: (1).
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NAME/KEY: misc_feature
                         FEATURE:
                                           OTHER INFORMATION: Description of Artificial Sequence: FIV9 primer
                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 15.7%;
Local Similarity 57.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45 CTTGGGGCCTTCTCGGCGAATTGGTTCACAGGACTTTGGTTTCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 CTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAA 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CTTGGGGCCTTCTCGGCGAATTGGTTCACAGGACTTTGGTTTCCA 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26; Conservative
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US-09-863-806-135; Sequence 135, Ap.; Publication No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Tang, Hengli
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PRODUCTION
TITLE OF INVENTION: OF RNA VIRUSES AND RNA VIRUS-BASED VECTOR PARTICLES
FILE REFERENCE: 1994-002001
CURRENT APPLICATION NUMBER: US/09/853,745
CURRENT FILING DATE: 2001-05-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/206,997
PRIOR FILING DATE: 2000-05-24
NUMBER OF SEQ ID NOS: 44
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Feng, Yu
APPLICANT: Tang, He
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (29)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: synthetically generated primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ocal Similarity
                                                                 PRIOR APPLICATION DATA:
                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 195
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: DETECTION OF NEOPLASIM BY ANALYSIS OF SALIVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sidransky, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            48 CATTATGTCTTTGTCACCCTGCTTTTGCTCCTATAGT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 CTCCATTCTGCCTGGGTGACAATGCCTTGG 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13 CCCCATCCTACCTTGTNGACTGTCCCTCGG 42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 CATTCTGCCTGGGTGACAATGCCTTGGTTTCAACACT 52
                                                                                     APPLICATION NUMBER: US/09/863,806 FILING DATE: 22-May-2001
                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FASTERS for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                              STREET: 4225 Executive Square, Suite 1400 CITY: La Jolla STATE: CA
                                                                                                                                                                                                                                                                                          COUNTRY:
                        APPLICATION NUMBER: 09/038,637 FILING DATE: <Unknown>
APPLICATION NUMBER: 08/152,313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09863806
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                                                                                                                                                                                                                                                                                          USA
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 Mismatches

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US-10-215-112-2201
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US-09-940-185-583/c
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                                                                                                                                                 Sequence 2201, Application US/10215112 Publication No. US20030082596A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/227, 948
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 4768
SOFTWARE: PatentIn version 3.1
SEQ ID NO 583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 583, Application US/09940185 Publication No. US20030096239A1 GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
                                                                                 TITLE OF INVENTION: Method of Genetic Analysis of Probes: TITLE OF INVENTION: Test3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/940,185
CURRENT FILING DATE: 2001-08-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Probes and Decoder Oligonucleotides FILE REFERENCE: A-69605-1
                                                                   FILE REFERENCE: 3119
                                                                                                                                APPLICANT: Michael Mittmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gunderson, Kevin APPLICANT: Chee, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Computer Generated Probe Sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 135: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 619/678-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE DESCRIPTION: SEQ ID NO: 135:
                                                                                                                                                                                                                                                                                                                                                 38 CTTGGTTTCAACACTATCGGAATG 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22 GCCTGGGTGACAATGC 37
                                                                                                                                                                                                                                                                                                         24 CTGGATTTCAACTCGCGCGGAATG 1
                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Haile, Lisa A. REGISTRATION NUMBER: 38,347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 12-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 07265/146001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 619/678-5099
                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                         15.5%; Score 14.4; DB 9; 75.0%; Pred. No. 4.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15.5%; Score 14.4; DB 9; Length 20; 93.8%; Pred. No. 4e+04;
                                                                                                                                                                                                                                                                                                                                                                                             0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 9; Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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; OTHER INFORMATION: Computer Generated Probe Sequence. US-09-940-185-4551
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4551, Application US/09940185; Publication No. US20030096239A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: Fast-SEQ for Windows Version 4.0
SEQ ID NO 13270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-940-185-4551/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Synthetic Oligonucleotide US-10-215-112-13270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   : GENERAL INFORMATION:

APPLICANT: Michael Mittmann

TITLE OF INVENTION: Method of Genetic Analysis of Probes:

TITLE OF INVENTION: Test3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 57
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; Publication No. US20030082596A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-215-112-13270/c
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                                                                                                                            SEQ ID NO 4551
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                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/227,948
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 4768
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/940,185
CURRENT FILING DATE: 2001-08-27
                                                                                                                                                                                                                                                                                                                                             APPLICANT: Gunderson, Kevin
APPLICANT: Chee, Mark
TITLE OF INVENTION: Probes and Decoder Oligonucleotides
                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: A-69605-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 3119
                                          FEATURE:
                                                             ORGANISM: Artificial Sequence
                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                       LENGTH:
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Local Similarity 75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28 GTGACAATGCCTTGGTTTCAACAC 51
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 Mismatches

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LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-098-263B-97583
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 97583, Application US/10098263B Publication No. US20030104410A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 5065, Application US/09866108 Patent No. US20020048800A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION UNMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: GU, Yizhong APPLICANT: JI, Yonggar APPLICANT: PENN, Shar:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/098,263B CURRENT FILING DATE: 2003-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Mittman, Michael TITLE OF INVENTION: Human Microarray FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/236,359 PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/866,108 CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: SHANNON, MATK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: ABONICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 131066
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PRIOR FILING DATE: 2001-03-16
     PRIOR APPLICATION NUMBER: PCT/US01/00663
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                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
                                                                                                                      APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
                         APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
                                                                        APPLICATION NUMBER: PCT/US01/00665 FILING DATE: 2001-01-30
                                                                                                                                                                     APPLICATION NUMBER: PCT/US01/00664 FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         63 GACTGTCGGGCCTCCAGAAAACAT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25 CTGGATTTCAACTCGCGCGGAATG 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 GACCCTCGAGACTCCAGAGGACAT 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JI, Yonggang
PENN, Sharron G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HANZEL, David K.
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75.0%;
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75.0%;
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Pred. No. 4.3e+04;
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US-09-866-108-5066/c
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; Patent No. US20020048800A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-866-108-5065
                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 6/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR PRILICATION NUMBER: PCT/US01/00667
PRIOR PRILICATION NUMBER: PCT/US01/00667
PRIOR PPLICATION NUMBER: PCT/US01/00664
PRIOR PPLICATION NUMBER: PCT/US01/00669
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR PRILING DATE: 2001-01-30
PRIOR PRILICATION NUMBER: PCT/US01/00668
PRIOR PRILICATION NUMBER: PCT/US01/00669
PRIOR PRILING DATE: 2001-01-30
PRIOR PRILICATION NUMBER: PCT/US01/00668
PRIOR PRILICATION NUMBER: PCT/US01/00668
PRIOR PRILICATION NUMBER: PCT/US01/00668
PRIOR PRILICATION NUMBER: PCT/US01/00668
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR PRILING DATE: 2001-01-30
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SEQ ID NO 5065
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                    PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT, PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: GU, Yizhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
                                                                   PRIOR APPLICATION NUMBER: PCT/US01/00670 PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: US 60/266,860
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                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/US01/00662
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                                              PRIOR APPLICATION NUMBER: US 60/234,687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                                                                                                                                           PCT/US01/00661
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; OTHER INFORMATION: Description of Artificial Sequence:primer BCD2N US-09-126-816-2
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NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 5066
                                                                                                                            SEQ ID NO 3
                                                                                                                                                                                                                                                                                         APPLICANT: BOEHRINGER MANNHEIM GMBH
TITLE OF INVENTION: METHOD OF INACTIVATION OF P21 RAS AND AGENTS THEREFOR
FILE REFERENCE: VON EICHEL-STREIBER
CURRENT APPLICATION NUMBER: US/09/126, 816
CURRENT FILING DATE: 1998-07-31
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                                                                                                                                           EARLIER APPLICATION NUMBER: PCT/EP97/00426
EARLIER FILING DATE: 1997-01-31
EARLIER APPLICATION NUMBER: 96 101 469.3
EARLIER FILING DATE: 1996-02-02
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
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EARLIER FILING DATE: 1997-01-31
EARLIER APPLICATION NUMBER: 96 101 469.3
EARLIER FILING DATE: 1996-02-02
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OTHER INFORMATION: Description of Artificial Sequence:primer BCD2C
                               FEATURE:
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                                                   ORGANISM: Artificial Sequence
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; GENERAL INFORMATION:
                                                                                           US-10-017-178-10; Sequence 10, Application US/10017178
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                                                                                ; Patent No. US200
                                                                                                                                         RESULT 65
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; Patent No. US20020006407A1
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TITLE OF INVENTION: High Throughput Assay to Detect Inhibitors of the MAP Kinase P
                     APPLICANT: Yamamoto, Hirotaka APPLICANT: Moskal, Joseph R.
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APPLICANT: O'Brien, Alison D.
APPLICANT: Wachtel, Marian R.
TITLE OF INVENTION: Histidine.
TITLE OF INVENTION: Maning Intin
TITLE OF INVENTION: Antigen Car
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REFERENCE/DOCKET NUMBER: 04
TELECOMMUNICATION INFORMATION:
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CURRENT APPLICATION DATA:
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Local Similarity 75.0%;
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1300 I Street, N.W., Suite 700
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N: 530
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                                                                                                                                                                                                                                                                                              15.5%;
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; Sequence 35, Application US/09795006A
; Patent No. US20020151680A1
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                                                                                                     RESULT 67
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                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/971,372A CURRENT FILING DATE: 2001-10-04 PRIOR APPLICATION NUMBER: PCT/US00/09021 PRIOR FILING DATE: 2000-04-05 PRIOR APPLICATION NUMBER: 60/127,912 PRIOR FILING DATE: 1999-04-06
APPLICANT: Alitalo et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Recombinant influenza viruses for vaccines and gene TITLE OF INVENTION: therapy FILE REFERENCE: 960296.98130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Kawaoka, Yoshihiro APPLICANT: Neumann, Gabriele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/017,178 CURRENT FILING DATE: 2001-12-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/255,548 PRIOR FILING DATE: 2000-12-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 99,123-D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: BsmBI recognition sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature LOCATION: (7)..(12)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artifical Sequence: PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Influenza virus sequence
                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature LOCATION: (17)..(46)
                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 46
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                                                                                                                                                                                                                                                            ocal Similarity
                                                                                                                                                                                             15 CCATTCTGCCTGGGTGACAATGCCTTGGTTTC 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 TCTGCCTGGGTGACAATGCCTTGG 42
                                                                                                                                                         46 CCATTATGTCTTTGTCACCCTGCTTTTGCTCC 15
                                                                                                                                                                                                                                         21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18; Conservative
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                                                                                                                                                                                                                                                        15.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gabriele
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75.0%;

 Mismatches

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                                                                                                                                                                                                                                                        Score 14.4; DB 9;
Pred. No. 5.2e+04;
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Pred. No. 4.8e+04,
                                                                                                                                                                                                                                                                             DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 12;
                                                                                                                                                                                                                                                                           Length 46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 35;
                                                                                                                                                                                                                                      Indels
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US-10-078-337-26/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 26, Application US/10078337 Publication No. US20030023068A1
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NUMBER OF SEQ ID NOS: 175
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CURRENT APPLICATION NUMBER: US/09/795,006A
CURRENT FILING DATE: 2001-02-26
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                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Artificial chimeric
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                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: STEFFE, ERIC K.
REGISTRATION NUMBER: 36,688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                            TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENDRESS, GREGORY A.
TITLE OF INVENTION: NOVEL HUMAN GROWTH FACTORS
NUMBER OF SEQUENCES: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: YU,
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                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/10/078,337 FILING DATE: 21-Feb-2002
                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                               TELEFAX: (202)
                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/916,576 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
TOPOLOGY: linear
                                                                                                                                               TELEPHONE: (202) 371-2600
TELEFAX: (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
                                                                      ENGTH: 49 base pairs
                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 1488.0500001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DILLON, PATRICK EBNER, REINHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1100 NEW YORK AVENUE, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GUO-LIANG
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75.0%;
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                                                                                                                       US-09-864-636A-2543
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-060-301-175/c
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                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/864,636A CURRENT FILING DATE: 2002-10-15 NUMBER OF SEQ ID NOS: 2640 SOFTWARE: Patentin version 3.0 SEQ ID NO 2543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: JP 2001-25700
PRIOR FILING DATE: 2001-02-01
NUMBER OF SEQ ID NOS: 200
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 175
                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Third wave Technologies
APPLICANT: Aliwal Hatim
APPLICANT: Bartholomay, Christian
APPLICANT: Chehak, Luanne
TITLE OF INVENTION: Detection of RNA Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: NAKAMURA, YUSUKe et al.
TITLE OF INVENTION: A METHOD FOR SNP (SINGLE NUCLEOTIDE POLYMORPHISM) TYPING
FILE REFERENCE: 1254-0195p
                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: FORS-04944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/060,301 CURRENT FILING DATE: 2002-02-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                      OTHER INFORMATION: Synthetic
                                                                                                                                                            FEATURE:
                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                       ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
                                                        Match 15.3%; Score 14.2; DB 9; Length 20; Local Similarity 84.2%; Pred. No. 4.8e+04;
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61 GTGACTGTCGGGCCTCCAG 79
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Conservative
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Similarity 84.2%;
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Similarity 60.0%;
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Pred. No. 4.8e+04;
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                                                                                                                 : Sequence 78553, Application US/10098263B 
: Publication No. US20030104410A1 
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 20064, Application US/10098263B; Publication No. US20030104410A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 71
US-10-098-263B-20064/c
                                                                                                                                                                               US-10-098-263B-78553
                                                                                                                                                                                                  RESULT 73
                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-098-263B-50784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 50784, Application US/10098263B; Publication No. US20030104410A1
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PRIOR FILING DATE: 2001-03-16
I NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 20064
LENGTH: 25
TYPER -----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 50784
                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                      APPLICANT: Mittman, Michael TITLE OF INVENTION: Human MFILE REFERENCE: 3118.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
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TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
                 CURRENT APPLICATION NUMBER: US/10/098,263B CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapien
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                                                                                                                                                                                                                                  Match 15.3%;
Local Similarity 84.2%;
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                                                                                                                                                                                                                                                                25 ACAATATCTGAATGTGAAT 7
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                                                                          Human Microarray
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Pred. No. 5.2e+04;
                                                                                                                                                                                                                                                                                                                                                                  Score 14.2; DB 9; Length 25; Pred. No. 5.2e+04;
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                                                                                                                                                                                                                                                                                                                                                Mismatches
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; Sequence 22, Application US/09872349
; Patent No. US20020132980A1
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APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric S.
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Sidney Pestka
TITLE OF INVENTION: PHOSPHORYLATED PROTEINS AND USES RELATED THERETO
FILE REFERENCE: PBLI-P01-007
CURRENT APPLICATION NUMBER: 05/09/872,349
CURRENT FILING DATE: 2001-05-31
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                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/255,296
PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 46
                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/208,240 PRIOR FILING DATE: 2000-05-31
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PRIOR FILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
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CURRENT FILING DATE: 2001-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
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OTHER INFORMATION: Description of Artificial Sequence: primer
                                                                                 ORGANISM: Artificial Sequence
                                                        FEATURE:
                                                                                                                                 TYPE: DNA
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                                                                                                                                                                  ENGTH:
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US-09-829-124-5/c
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                                                                                                                                                                                          SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                        FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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Sequence 5, Application US/09829124
Patent No. US20020066122A1
                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/829,124
CURRENT FILING DATE: 2001-04-09
PRIOR APPLICATION NUMBER: 60/224,053
PRIOR FILING DATE: 2000-08-09
PRIOR APPLICATION NUMBER: 09/412,452
                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: HYPERSENSITVE RESPONSE ELICITOR FROM XANTHOMAS TITLE OF INVENTION. CAMPESTRIS FILE REFERENCE: 21629/L01
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CURRENT FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/256,349
PRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wei, Zhong-Min
APPLICANT: Swanson, Shana
APPLICANT: Fan, Hao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: STAPHYLOG
TITLE OF INVENTION: STAAU_R9
FILE REFERENCE: 073406-0701
                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/103,124 PRIOR FILING DATE: 1998-10-05
                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1999-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: COMPOSITIONS AND METHODS INVOLVING AN ESSENTIAL TITLE OF INVENTION: STAPHYLOCOCCUS AUREUS GENE AND ITS ENCODED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: PELLETIER, JERRY APPLICANT: GROS, PHILIPPE APPLICANT: DUBOW, MICHAEL
OTHER INFORMATION: Description of Artificial Sequence: primer
                                          TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 15.3%; Score 14.2; DB 9; Length 33; Local Similarity 70.4%; Pred. No. 5.7e+04; es 19; Conservative 0. Nicolary 15.7e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             34 ATGCCTTGGTTTCAACACTATCGGAAT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32 GACCCAGGACCACTCCATGGTGGCTCG 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GAGCTGGGAGCACTCCATTCTGCCTGG 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19; Conservative
                                                                                                                                                              PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PatentIn Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GROS, PHILIPPE
DUBOW, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fan, Hao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Swanson, Shane S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15.3%;
70.4%;
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                                                     US-08-781-986A-2061/c
                                                                         RESULT 79
                                                                                                                                                                                                                                                                                   US-10-043-573-131
                  Sequence 2061, Application US/08781986A Publication No. US20030054436A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 131, Applicat Publication No. US2003 GENERAL INFORMATION
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                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                             Query Match
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 131:
                                                                                                                                                                                              15.3%;
Local Similarity 65.5%;
es 19; Conservation
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FILING DATE: 07-MAR-1997
APPLICATION NUMBER: US 60/032,069
FILING DATE: 02-DEC-1996
ATTORNEY/AGENT INFORMATION:
NAME: Liebeschuetz, Joe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/043,573
FILING DATE: 09-Jan-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 018547-030100US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: <Unknown> PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sapolsky, Ronald J. TITLE OF INVENTION: Brassica Polymorphisms NUMBER OF SEQUENCES: 173
                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 131:
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lemieux, Bertrand
Landry, Benoit S.
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                                                                                                                                                                   42 GTTTCAACACTATCGGAATGTGACTGTCG 70
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                                                                                                                                4 GTATCAGGACAATGTGTKWGTGACTGTGG 32
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                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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STATE: California
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                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 42 base pairs
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    Mismatches

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                                                                                                                                                                                                                                             Score 14.2; DB 9; Length 42;
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US-09-940-185-517/c
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                                                                                                                                                                   SOFTWARE: Pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                              FEATURE:
                                                                                                                                               LENGTH: 24
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CTTY: Rockville
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16;
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; OTHER INFORMATION: Computer Generated Probe Sequence. US-09-940-185-517
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/227,948
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/228,854
                                                                                                                                                                                                             PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 4768
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/940,185
CURRENT FILING DATE: 2001-08-27
                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Probes and Decoder Oligonucleotides FILE REFERENCE: A-69605-1
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Gunderson, Kevin APPLICANT: Chee, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
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LENGTH: 50 base pairs
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MEDIUM TYPE: Diskett
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Charles Kunsch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: MSI
OPERATING SYSTEM: MSI
OPERATING SYSTEM: ASCII Text
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/781,986A
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Local Similarity 84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41 GGTGACAATATCTTTGTTT 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27 GGTGACAATGCCTTGGTTT 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9410 Key West Avenue
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(301) 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Diskette, 3.50 inch, 1.4Mb storage
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  15.1%; Score 14; DB 9; 77.3%; Pred. No. 6.1e+04;
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 Mismatches

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        US-10-215-112-5487
                               RESULT 83
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 919
                                                                                                                                                                      Matches
                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Publication No. US20030082596A1
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Publication No. US20030096239A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Method of Genetic Analysis of Probes: TITLE OF INVENTION: Test3
FILE REFERENCE: 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEG ID NOS: 4768
SOFTWARE: Patentin version 3.1
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CURRENT FILING DATE: 2001-08-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Michael Mittmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/227,948
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
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                                                                                                                                                                                          Local Similarity
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                                                                                                          30 GACAATGCCTTGGTTTCAACAC 51
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                                                                                                                                                                      Conservative
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                                                                                                                                                                                      Score 14; DB 9;
Pred. No. 6.2e+04
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                                                                                                                                                                                                          DB 9; Length 25;
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US-10-098-263B-62496
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                                                                                                                        ; GENERAL INFORMATION:
                                                                                                                                       ; Sequence 62496, Application US/10098263B ; Publication No. US20030104410A1
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/098,263B CURRENT FILING DATE: 2003-01-08 PRIOR APPLICATION NUMBER: 60/276,759
                                                            APPLICANT: Mittman, Michael TITLE OF INVENTION: Human Microarray FILE REFERENCE: 3118.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2001-08-29
NUMBER OF SEG ID NOS: 4768
SOFTWARE: PatentIn version 3.1
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CURRENT FILING DATE: 2001-08-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Probes and Decoder Oligonucleotides FILE REFERENCE: A-69605-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gunderson, Kevin APPLICANT: Chee, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 3119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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PRIOR FILING DATE: 2001-03-16

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; Publication No. US20
; GENERAL INFORMATION:
; APPLICANT: Mittman,
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Publication No. US20030104410A1
GENERAL INFORMATION:
                                                                                                                                               US-10-098-263B-80753
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CURRENT FILING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 76584
LENGTH: 25
                                                               Matches
                                                                                                                                                                                                                         SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1 SEQ ID NO 80753
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.SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 62496
LENGTH: 25
                                                                                                      Query Match
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CURRENT FILLING DATE: 2003-01-08
PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
NUMBER OF SEQ ID NOS: 131066
COUNTAINE MATERIAL DEPORT OF SEQ ID NOS: 131066
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Mittman, Michael
TITLE OF INVENTION: Human Microarray
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TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
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                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapien
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                                                        Local Similarity es 17; Conserva
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                     55 CGGAATGTGACTGTCGGGCCTC 76
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Conservative
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77.3%;
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 Mismatches

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Pred. No. 6.2e+04;
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Pred. No. 6.2e+04,
                                                            Mismatches
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US-10-097-672-3/c
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                                                                                                                                         ; GENERAL INFORMATION:
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             APPLICANT: Boston College
APPLICANT: Braley, Andrew
APPLICANT: McLaughlin, Lar
APPLICANT: DongLi, Chen
APPLICANT: Lan, Tao
TITLE OF INVENTION: Selecti
                                                                                                                                                            Sequence 3, Application US/10097672 Publication No. US20030100759A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 112182
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TITLE OF INVENTION: Human Microarray
FILE REFERENCE: 3118.1
FILE REFERENCE: 2846/2012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/276,759 PRIOR FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Mittman, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/276,759 PRIOR FILING DATE: 2001-03-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 131066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Human Microarray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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                                                                         Fraley, Andrew W
McLaughlin, Larry W
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                   Selective Anti-Viral Nucleoside Chain Terminators
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Pred. No. 6.2e+04;
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Pred. No. 6.2e+04;
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US-09-912-263-506
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                                                                          GENERAL INFORMATION:
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                                                                                           Publication No. US20030039973A1
                                                                                                             Sequence 506,
                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 103
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APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric. S.
APPLICANT: Lander, Eric. S.
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CURRENT FILING DATE: 2001-07-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 09/145,828 PRIOR FILING DATE: 1998-09-02
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TITLE OF INVENTION: ELONGASE GENES AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Primer RO720
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                                                                                                                                                                                                                                                                                             Similarity
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Leonard, Amanda Eun-Yeong
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                                                                                                           Application US/09912263
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66.7%;
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66.7%;
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Pred. No.
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Pred. No. 6.6e+04;
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TYPE: DNA
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US-10-029-314-21; Sequence 21, Application US/10029314; Publication No. US20030077701A1
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; Sequence 20, Application US/10029314
; Publication No. US20030077701A1
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CURRENT APPLICATION NUMBER: US/10/029,314
CURRENT FILING DATE: 2001-12-20
PRIOR APPLICATION NUMBER: 09/899,732
PRIOR FILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: 09/610,635
PRIOR APPLICATION NUMBER: 09/610,635
PRIOR FILING DATE: 2000-07-05
NUMBER OF SEO ID NOS: 28
SOFTWARE: Patentin Ver. 2.1
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PRIOR FILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 28
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CURRENT FILING DATE: 2001-12-20
PRIOR APPLICATION NUMBER: 09/899,732
PRIOR FILING DATE: 2001-07-05
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CURRENT FILING DATE: 2001-07-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: primer/probe
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Sequence 20, Application US/09899732
Publication No. US20030082623A1
GENERAL INFORMATION:
APPLICANT: Salon et al, John A.
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US-09-899-732-20/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21, Application US/09899732 Publication No. US20030082623A1 GENERAL INFORMATION:
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LENGTH: 32
TYPE: DNA
                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 28
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/899,732
CURRENT FILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: 09/610,635
PRIOR FILING DATE: 2000-07-05
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 20
     Query Match
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                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/899,732
CURRENT FILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: 09/610,635
PRIOR FILING DATE: 2000-07-05
                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Salon et al. John A. TITLE OF INVENTION: DNA Encoding A Human Melanin Concentrating Hormone TITLE OF INVENTION: Receptor (MCHL) And Uses Thereof EILE REFERENCE: 1795/57453-C/JPW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: DNA Encoding A Human Melanin Concentrating Hormone TITLE OF INVENTION: Receptor (MCH1) And Uses Thereof FILE REFERENCE: 1795/57453-C/JPW
                                                                                                                                                 LENGTH: 32
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                       OTHER INFORMATION: Description of Artificial Sequence: primer/probe
                                                                                                    FEATURE:
                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      match 15.1%; Score 14; DB 9; Length 32; Local Similarity 66.7%; Pred. No. 6.7e+04; es 20; Conservative 0. Michael 1
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15.1%; Score 14; DB 9; Length 32;
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NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.1
SEQ ID NO 20
LENGTH: 32
TYPE: DNA
ORGANISM: ARTIFICIAL SEQUENCE
                                                                                      US-09-885-478-21
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US-09-885-478-21
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                                                                                                                                                                                                            SOFTWARE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21, Application U Patent No. US20020111306A1 GENERAL INFORMATION:
                                        Query Match
  Matches 20;
                        Best
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APPLICANT: LAZ, THOMAS M
APPLICANT: LAZ, THOMAS M
APPLICANT: NAGORNY, RAISA
APPLICANT: WILSON, AMY E
APPLICANT: WILSON, AMY E
TITLE OF INVENTION: DAE SHOODING A HUMAN MELANIN CONCENTRATING HORMONE RECEPTOR FILE REFERENCE: 1795/57453-A-PCT-US
CURRENT FALLING NUMBER: USES THEREOF
FILE REFERENCE: 1795/57453-A-PCT-US
CURRENT FILING DATE: 2001-09-24
PRIOR APPLICATION NUMBER: PCT/US99/31169
PRIOR APPLICATION NUMBER: PCT/US99/31169
PRIOR FILING DATE: 1999-12-30
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TITLE OF INVENTION: DNA ENCODING A HUMAN MELANIN CONCENTRATING HORMONE RECEPTOR (M
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 1795/57453-A-PCT-US
CURRENT APPLICATION NUMBER: US/09/885,478
CURRENT FILING DATE: 2001-09-24
PRIOR APPLICATION NUMBER: PCT/US99/31169
PRIOR FILING DATE: 1999-12-30
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                                                                                                                                 TYPE: DNA
ORGANISM: ARTIFICIAL SEQUENCE
                                                                                                          OTHER INFORMATION: PRIMER
                                                                                                                            FEATURE:
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                                                                                                                                                                                       LENGTH: 32
                  y Match 15.18;
Local Similarity 66.78;
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  Conservative
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                    Score 14; DB 10; Length 32; Pred. No. 6.7e+04;
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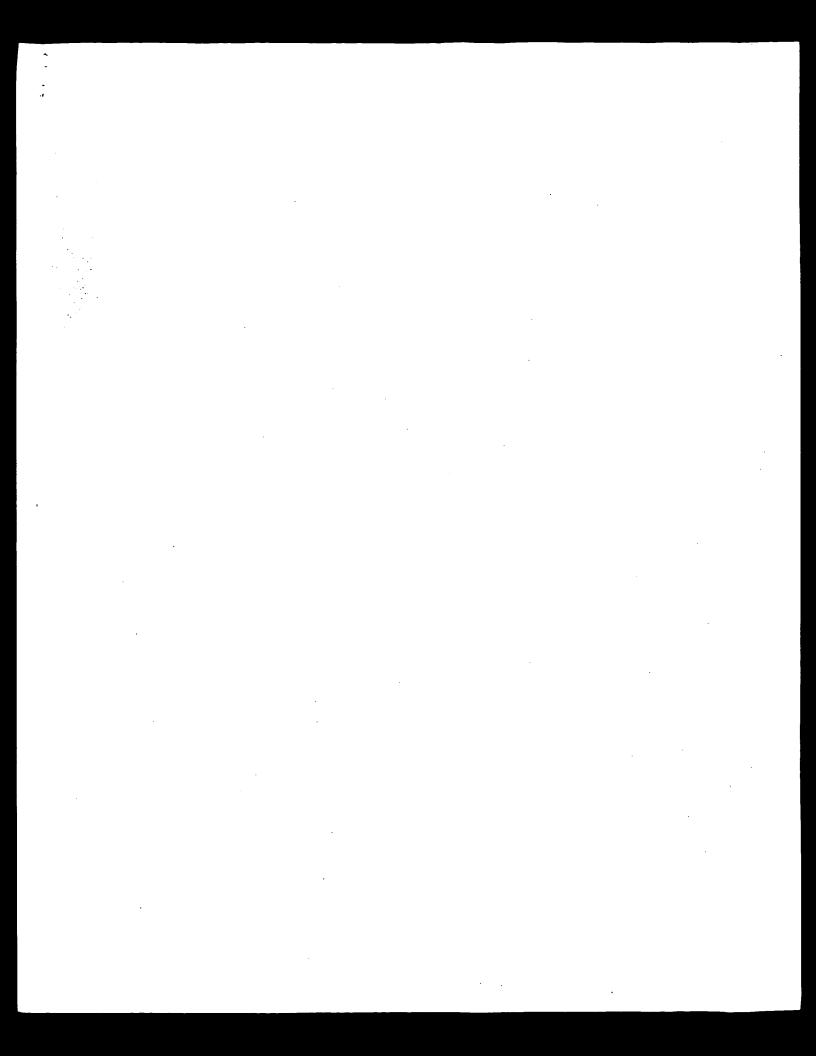
11 CACTCCATTCTGCCTGGGTGACAATGCCTT 40

Job time : 123 secs

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CURRENT APPLICATION NUMBER: US/10/005,956
CURRENT-FILING DATE: 2001-12-03
PRIOR APPLICATION_NUMBER: 60/251,015
PRIOR FILING DATE: 2000-12-04
PRIOR APPLICATION_NUMBER: 60/263,678
PRIOR FILING DATE: 2001-01-23
PRIOR FILING DATE: 2001-03-03
PRIOR FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 1579
SOFTWARE: PATENTIAN VERSION 3.0
SEQ ID NO 1436
LENGTH: 39
TYPEN: NMA
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                                                                                                                                                                                                                       ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-005-956-1436
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 99
US-10-102-949-2/c
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US-10-005-956-1436/c
Search completed: July 5, 2003, 13:11:58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1436, Application US/10005956 Publication No. US20030113726A1
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Patent No. US20020157140A1
                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ESRI - A PLANT GENE THAT CAN PROMOTE PLANT REGENERATION
TITLE OF INVENTION: AND TRANSFORMATION
FILE REFERENCE: 2312-107-2
CURRENT APPLICATION NUMBER: US/10/102,949
CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: US/09/604,394
PRIOR PELLING DATE: 2000-06-27
NUMBER: 05 SEQ ID NOS: 13
SOFTWARE: PATENTIN Ver. 2.0
                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: D0053NP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Banno, Hiroharu
APPLICANT: Chua, Nam-Hai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                        Local Similarity 66.7 es 20; Conservative
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                                                                                                     18 TTCTGCCTGGGTGACAATGCCTTGGTTTCA 47
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66.78;
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 Mismatches

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Post-processing: Minimum Match 0%
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Maximum DB seq length: 50
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            /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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971 12.2 13.1 20 1 US-08-715-142-5 Sequence 972 12.2 13.1 20 3 US-08-281-203-2 Sequence 973 12.2 13.1 20 3 US-08-79-764A-45 Sequence 974 12.2 13.1 20 4 US-09-180-437-50 Sequence 975 12.2 13.1 20 4 US-09-180-437-51 Sequence	970 12.2 13.1 19 4 US-19-559-565C-68 Sequence 969 12.2 13.1 19 4 US-9-559-565C-69 Sequence 970 12.2 13.1 20 1 US-08-564-002-2 Sequence	966 12.2 13.1 19 4 US-09-344-002B-68 Sequence 967 12.2 13.1 19 4 US-09-344-002B-69 Sequence	964 12.2 13.1 19 4 US-08-769-062B-68 Sequence 965 12.2 13.1 19 4 US-08-769-062B-69 Sequence	963 12.2 13.1 19 4 US-09-339-904A-69 Sequence	c 961 12.2 13.1 19 4 US-09-339-913B-69 Sequence	c 959	c 958 12.2 13.1 19 1 US-08-501-626-50 Sequence	c 956 12.2 13.1 19 1 US-08-378-860-50 Sequence	c 955 12.2 13.1 19 1 US-08-501-779-50 Sequence	c 954 12.2 13.1 19 1 US-08-398-945-50 Sequence	c 952 12.2 13.1 18 4 US-08-869-423-8 Sequence	950 12.2 13.1 18 4 US-09-018-584A-146 Sequence	949 12.2 13.1 18 4 US-09-521-109-5 Sequence	c 947 12.2 13.1 18 3 US-09-344-520-12 Sequence	946 12.2 13.1 17 1 US-08-717-526-42 Sequence	c 944 12.4 13.3 50 5 PCT-US95-04632-6 Sequence c 945 12.4 13.3 50 5 PCT-US96-07627-17 Seguence	942 12.4 13.3 50 4 US-09-261-115-16 Sequence c 943 12.4 13.3 50 5 PCT-US93-12388-406 Sequence	c 941 12.4 13.3 50 4 US-08-889-56/A-7/ Sequence c 941 12.4 13.3 50 4 US-09-354-947-406 Sequence	939 12.4 13.3. 50 4 US-09-194-301-1 Sequence	c 937 12.4 13.3 50 3 US-08-937-195-17 Sequence c 938 12.4 13.3 50 4 US-08-915-152-17 Sequence	c 935 12.4 13.3 50 2 US-08-475-228A-406 Sequence c 936 12.4 13.3 50 3 US-08-482-080A-406 Sequence	c 933 12.4 13.3 50 1 US-08-231-227-6 Sequence c 934 12.4 13.3 50 1 US-08-123-936-406 Sequence	c 932 12.4 13.3 49 4 US-09-380-190A-37 Sequence c 932 12.4 13.3 50 1 US-08-171-389-406 Sequence	930 12.4 13.3 49 4 US-09-813-781-69 Sequence	928 12.4 13.3 48 4 US-09-548-260-50 Sequence	927 12.4 13.3 48 4 US-09-390-867A-50 Sequence	925 12.4 13.3 47 5 PCT-US94-14436-13 Sequence	c 923 12.4 13.3 47 4 US-09-641-638-778 Sequence	c 921 12.4 13.3 47 4 US-09-218-207-258 Sequence 922 12.4 13.3 47 4 US-09-641-638-777 Sequence	920 12.4 13.3 47 4 US-09-338-907-258 Sequence	c 918 12.4 13.3 47 3 US-08-155-938-1 Sequence	915 12.4 13.3 46 3 US-09-249-215-107 Sequence	915 12.4 13.3 46 2 US-08-910-408-107 Sequence	913 12.4 13.3 45 5 PCT-US94-06197-4 Sequence	912 12.4 13.3 45 5 PCT-US93-12388-222 Sequence	910 12.4 13.3 45 4 US-09-354-947-222 Sequence	908 12.4 13.3 45 4 US-08-639-294-4 Sequence	906 12.4 13.3 45 3 US-08-482-080A-222 Sequence 907 12.4 13.3 45 4 US-08-987-943-11 Seguence	0006 12 4 13 3 45 105-08-482-080A-62 Sequence
971 12.2 13.1 20 1 972 12.2 13.1 20 3 973 12.2 13.1 20 3 974 12.2 13.1 20 4 975 12.2 13.1 20 4	970 12.2 13.1 19 4 US-19-559-565C-68 Sequence 969 12.2 13.1 19 4 US-9-559-565C-69 Sequence 970 12.2 13.1 20 1 US-08-564-002-2 Sequence	966 12.2 13.1 19 4 US-09-344-002B-68 Sequence 68 967 12.2 13.1 19 4 US-09-344-002B-69 Sequence 69	964 12.2 13.1 19 4 US-08-769-062B-68 Sequence 965 12.2 13.1 19 4 US-08-769-062B-69 Sequence	963 12.2 13.1 19 4 US-09-339-904A-69 Sequence	c 961 12.2 13.1 19 4 US-09-339-913B-69 Sequence	c 959	c 958 12.2 13.1 19 1 US-08-501-626-50 Sequence	c 956 12.2 13.1 19 1 US-08-378-860-50 Sequence	c 955 12.2 13.1 19 1 US-08-501-779-50 Sequence	c 954 12.2 13.1 19 1 US-08-398-945-50 Sequence	c 952 12.2 13.1 18 4 US-08-869-423-8 Sequence	950 12.2 13.1 18 4 US-09-018-584A-146 Sequence	949 12.2 13.1 18 4 US-09-521-109-5 Sequence	c 947 12.2 13.1 18 3 US-09-344-520-12 Sequence	946 12.2 13.1 17 1 US-08-717-526-42 Sequence	c 944 12.4 13.3 50 5 PCT-US95-04632-6 Sequence c 945 12.4 13.3 50 5 PCT-US96-07627-17 Seguence	942 12.4 13.3 50 4 US-09-261-115-16 Sequence c 943 12.4 13.3 50 5 PCT-US93-12388-406 Sequence	C 941 12.4 13.3 50 4 US-08-849-56/A-7/	939 12.4 13.3. 50 4 US-09-194-301-1 Sequence	c 937 12.4 13.3 50 3 US-08-937-195-17 Sequence c 938 12.4 13.3 50 4 US-08-915-152-17 Sequence	c 935 12.4 13.3 50 2 US-08-475-228A-406 Sequence c 936 12.4 13.3 50 3 US-08-482-080A-406 Sequence	c 933 12.4 13.3 50 1 US-08-231-227-6 Sequence c 934 12.4 13.3 50 1 US-08-123-936-406 Sequence	c 932 12.4 13.3 49 4 US-09-380-190A-37 Sequence c 932 12.4 13.3 50 1 US-08-171-389-406 Sequence	930 12.4 13.3 49 4 US-09-813-781-69 Sequence	928 12.4 13.3 48 4 US-09-548-260-50 Sequence	927 12.4 13.3 48 4 US-09-390-867A-50 Sequence	925 12.4 13.3 47 5 PCT-US94-14436-13 Sequence	c 923 12.4 13.3 47 4 US-09-641-638-778 Sequence	c 921 12.4 13.3 47 4 US-09-218-207-258 Sequence 922 12.4 13.3 47 4 US-09-641-638-777 Sequence	920 12.4 13.3 47 4 US-09-338-907-258 Sequence	c 918 12.4 13.3 47 3 US-08-155-938-1	915 12.4 13.3 46 3 US-09-249-215-107 Sequence	915 12.4 13.3 46 2 US-08-910-408-107 Sequence	913 12.4 13.3 45 5 PCT-US94-06197-4 Sequence	912 12.4 13.3 45 5 PCT-US93-12388-222 Sequence	910 12.4 13.3 45 4 US-09-354-947-222 Sequence	908 12.4 13.3 45 4 US-08-639-294-4 Sequence	906 12.4 13.3 45 3 US-08-482-080A-222 Sequence 223 907 12.4 13.3 45 4 US-08-987-943-11 Sequence 11.	000 12.4 13.3 45 3 US-08-48Z-08UA-6Z Sequence 62

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US-08-602-716A-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                 INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
          SEQUENCE CHARACTERISTICS:
LENGTH: 45 base pairs
                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/060,560
FILING DATE: 13-MAY-1993
                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: BASHAM, Daryl A.
APPLICANT: MILLER, Jeanette C.
TITLE OF INVENTION: PSYCHOSIS PROTECTING NUCLEIC ACID,
TITLE OF INVENTION: PEPTIDES, COMPOSITIONS AND METHODS OF USE
                                                                TELECOMMUNICATION INFORMATION:
                                                                                                 APPLICATION NUMBER: PCT/U
FILING DATE: 13-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, Roger L.
                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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                                 TELEPHONE: 202-737-3528
                                                                          NAME: BROWDY, ROGER L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: FR
                                                                                                                                                                                                 APPLICATION NUMBER: US/OFFILING DATE: 23-FEB-1996
TYPE: nucleic acid
                                                                                                                                                                                        CLASSIFICATION: 435
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US-08-446-033-48
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US-09-387-341-204
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Sequence 204, App
Sequence 53, Appl
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Sequence 75,
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Sequence 48,
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; TOPOLOGY: linear
; MOLECULE TYPE: CDNA
US-08-602-716A-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 2
US-08-331-081B-14
                                                                                                                                                                                                                        Matches
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            GENERAL INFORMATION:
APPLICANT: PASTAN
                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (613) 234-5233 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Devlin, Robert H. TITLE OF INVENTION: Transger
APPLICANT: KREITMAN, Robert J.
                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: Canal FILING DATE: June 17, 19 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                            TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 26, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: Octobe
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OPERATING SYSTEM: PC Dos 5.0
SOFTWARE: Wordperfect 5.1 (Patseq.txt)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 225 I
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STREET: 225 Metcalfe Street, Suite 606
                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
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                                                                                                                                                                                                                        Conservative
                  PASTAN, Ira
                                                                                                                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                                      17.8%; Score 16.6; DB 2; Length 30; 82.6%; Pred. No. 6.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Canada 2,126,138
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Pred. No. 1.2e+02;
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; MOLECULE TYPE: DNA (primer) US-08-225-224-31
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INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Pastan, Ira
APPLICANT: Kreitman, Robert J.
APPLICANT: Puri, Raj K.
                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Circularly Permuted Ligands and TITLE OF INVENTION: Circularly Permuted Chimeric Molecules NUMBER OF SEQUENCES: 72
                       PRIOR APPLICATION DATA:
                                                                                                                                                                                               COMPUTER READABLE FORM:
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COMPUTER READABLE FORM:
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                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Palannament
  APPLICATION NUMBER:
                                                                                                                                                                                                                                                         STREET: 'rwo .....
CITY: San Francisco
City: San Francisco
                                         CLASSIFICATION:
                                                          APPLICATION NUMBER: US/08/722,258 FILING DATE: 08-JAN-1997
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FILING DATE: 8-APR-1994
CLASSIFICATION: 530
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REGISTRATION NUMBER: 32,762
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Local Similarity 71.0%;
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Steuart Street Tower, One Market Plaza
                                                                                                                                                                                                                                                                                                 Two Embarcadero Center, Eighth Floor
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                                                                                                                                                                                                                                            USA
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                                                                                                                                                                                                                                                                                                                  Townsend and Townsend and Crew LLP
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WO PCT/US95/04468
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PCT-US95-04468-31
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   Best Local Similarity Matches 22; Conserv
                                      Query Match
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INFORMATION FOR SEQ ID NO: (
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                                                                                     MOLECULE TYPE: DNA (primer)
                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,7
                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200
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                                                                                                          TOPOLOGY:
                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/225,224
                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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OTHER INFORMATION:
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REGISTRATION NUMBER: 32,7
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                                                                                                                                        nucleic acid
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   Conservative
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15) 576-0300
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71.0%;
17.8%; Score 16.6; DB 5; 71.0%; Pred. No. 7.3e+02; tive 0; Mismatches 9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16.6; DB 3; Length 39; Pred. No. 7.3e+02;
                                    DB 5; Length 39;
   Indels
 0;
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 Gaps
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43 TTTCAACACTATCGGAATGTGACTGTCGGGC 73

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RESULT 7
US-08-384-708A-5
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                                                                                Sequence 5, Application US/08384708A Patent No. 5639868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence 34, Application US/08198670A Patent No. 5707796
                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                       Query Match
                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX 04/C
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 49 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 07/53
FILING DATE: 11-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NAME: Barry J. Swanson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: GOLD, LAKKY
APPLICANT: BEUTEL, BRUCE
METH
APPLICANT: Gold, Larry
APPLICANT: Janjic, Nebojsa
TITLE OF INVENTION: High-Affinity RNA Ligands of Basic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 07/71 FILING DATE: 10-JUNE-1991 PRIOR APPLICATION DATA:
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On NIMBER: 07/960,093
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: METHOD FOR SELECTING NUCLEIC ACIDS TITLE OF INVENTION: ON THE BASIS OF STRUCTURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/198,670A FILING DATE: 22-FEBRUARY-1994
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                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 07/960,00 FILING DATE: 14-OCTOBER-1992
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                                                                                                                                                                                                                                                                                                      Match 17.8%; Score 16.6; DB 1; Length 49;
Local Similarity 71.0%; Pred. No. 8.1e+02;
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                                                                                                                                                                                                    19 CACTATAGGGAGATGCCTGTCGAGCATGCTG 49
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Englewood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (303) 793-3333
303) 793-3433
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 Mismatches

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; Patent No. 6177557
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                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                               APPLICANT: Gold, Larry
APPLICANT: Janjic, Nebojsa
APPLICANT: Tasset, Diane
TITLE OF INVENTION: HIGH-AF
TITLE OF INVENTION: FIBROBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 07/714,131 FILING DATE: 10-JUNE-1991 PRIOR APPLICATION DATA:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MB storage
                                                                                                                                            NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                    TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Barry J. Swanson REGISTRATION NUMBER: 33 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY:
                                                                          STATE:
                                                                                                         STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
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                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        49 CACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                                     20 CACTATAGGGAGATGCCTGTCGAGCATGCTG 50
                                       8011
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: Colorado
                                                                                          Englewood
                                                                        Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        50 base pairs
                                                                                                           8400 E. Prentice Avenue, Suite 200
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                                                        USA
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                                                                                                                             Swanson & Bratschun, L.L.C.
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71.0%;
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                                                                                                                                                                                    THROMBIN
                                                                                                                                                                                                   HIGH-AFFINITY LIGANDS OF BASIC FIBROBLAST GROWTH FACTOR AND
                                                                                                                                                                  445
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 Mismatches

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US-08-687-421-5
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                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Williams
APPLICANT: Madaio,
APPLICANT: Weiner,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (303) 793-34
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 07/536,428
FILING DATE: 11-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NAME: BAITY J. SWANSON
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEXO7/PCT
                  SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 50 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
TELEFAX: (303) 793-3433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                                                                                                                                                                      STREET: One Liberty Place, 46th floor CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: MS-DOS
APPLICATION NUMBER: US/08/957,001B
                                                  OPERATING SYSTEM: windows
                                                                       COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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                                                                                                                         19103
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                                                                                                                                                     Pennsylvania
                                                                                                                                                                                                                                                                             Weiner, David B.
                                                                                                                                                                                                                                                                                                           Williams, William V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                        USA
                                                             E: Floppy disk
IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17.8%;
71.0%;
                                                                                                                                                                                                                                                                                            Michael
                                                                                                                                                                                                                                                            IMPROVED VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07/714,131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.6; DB 4; Length 50; Pred. No. 8.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
LENGTH:
                                                                                                                                        NAME:
                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                            FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE:
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Sequence 19, Application US/09496301
Patent No. 6248565
                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 215-568-3439 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 47 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/029,592
FILING DATE: 23-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Weiner, David B.
TITLE OF INVENTION: IMPROVED VACCINES
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                     TELEPHONE: 215-568-3439
                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: UP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 23-OCT-1997
                                                                                                                REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: UP
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: windows SOFTWARE: Wordperfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 23-OCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ocal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         55 CGGAATGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CAGRYTGYTGCAGCTGGACCTCCTGATAAGATTTAGAT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23; Conservative
                                                                                                                                                  DeLuca, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DeLuca, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pennsylvania
47 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             One Liberty Place, 46th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Williams, William V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Madaio, Michae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                 23-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
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60.5%;
                                                                                                                                                                                                                                 us 60/029,592
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                                                                                                                     UPN-3303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.4; DB 4; Length 47; Pred. No. 9.6e+02;
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RESULT 12
US-09-276-533A-10/c
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US-08-778-494B-103/c
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                                                                                                                                          Matches
                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 103, Application US/08778494B Patent No. 5962272
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                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Zhu, York
APPLICANT: Diachenko
APPLICANT: Siebert,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                       NAME: Pace, Doran R. BREGISTATION NUMBER: 38,261
REFERENCE/DOCKET NUMBER: CL-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (352) 375-8100
                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 40 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 03-JAN-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 2421 N.W. 41st Street, Suite A-1 CITY: Gainesville
                                                                                                                                                                                                                                         STRANDEDNESS: si
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 03-JAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/778,494B
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                       Local
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                                                               37 GCTGGCTGCCCCACTCTGCGTTGTTACCACTGCTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      55 CGGAATGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CAGRYTGYTGCAGCTGGACCTCCTGATAAGATTTAGAT 9
                                                                                                 3 GCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                      24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23;
                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                            nucleic acid
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                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhu, York
Diachenko, Luda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chenchik, Alex
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                        DNA (synthetic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-JAN-1997
                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                                            372-5800
                                                                                                                                                   17.4%; Score 16.2; DB 2; Length 40; 64.9%; Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                METHODS AND COMPOSITIONS FOR FULL-LENGTH CDNA CLONING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17.6%; Score 16.4; DB 4; Length 47; 60.5%; Pred. No. 9.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US 08/582,562
                                                                                                                                                                                                                                                                                                                                 103:
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                                                                                                                                      Mismatches
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; TYPE: DNA
; ORGANISM: oligonucleotide
US-09-237-712-6
                                                                Sequence 45, Application US/08813507; Patent No. 6114116; GENERAL INFORMATION:
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                                                                                                                                       US-08-813-507-45/c
                                                                                                                                                             RESULT 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; SOFTWARE: PatentIn Ver. 2.0 ; SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; GENERAL INFORMATION:
; APPLICANT: BROWN, WILLIAM C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-237-712-6/c
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SEQ ID NO 10
LENGTH: 39
                                                                                                                                                                                                                                                                                                                           Matches
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Patent No. 6180391
                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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APPLICANT: Sands, Arthur T.
APPLICANT: Sands, Arthur T.
TITLE OF INVENTION: VECTORS FOR GENE MUTAGENESIS AND GENE
TITLE OF INVENTION: DISCOVERY
FILE REFERENCE: 07705-0060-00000
CURRENT APPLICATION NUMBER: US/09/276,533A
CURRENT APPLICATION NUMBER: US/09/079,729
PRIOR APPLICATION NUMBER: US/09/079,729
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
PRIOR PILING DATE: 1998-04-14
PRIOR PILING DATE: 1998-04-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10, Application US/09276533A Patent No. 6436707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/237,712 CURRENT FILING DATE: 1999-01-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HIGHLY EFFICIENT CONTROLLED EXPRESSION OF EXOGENOUS TITLE OF INVENTION: GENES IN E. COLI FILE REFERENCE: A-518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EARLIER APPLICATION NUMBER: 60/072,794
EARLIER FILING DATE: 1998-01-28
NUMBER OF SEQ ID NOS: 98
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APPLICANT: Friedrich, Glenn A.
APPLICANT: Lilleberg, Stan
APPLICANT: Sands, Arthur T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/109,302 PRIOR FILING DATE: 1998-11-20
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                         Match 17.0%; Score 15.8; DB 4;
Local Similarity 74.1%; Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 17.0%;
Local Similarity 74.1%;
                                                                                                                                                                                                                        27 CACTCAGTTCGTACTGGTTGACAAGGC 1
                                                                                                                                                                                                                                                      11 CACTCCATTCTGCCTGGGTGACAATGC 37
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Landry, Benoit S.
Sapolsky, Ronald J.
                                       Lemieux,
                                                                                                                                                                                                                                                                                                                      Conservative
                                              Bertrand
                                                                                                                                                                                                                                                                                                                      0;

 Mismatches

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                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 40;
                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                      Gaps.
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US-08-813-507-45
                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-464-453-45/c
                                                                                                                                                                                                                                                                                                                                                            Patent No. 6358686
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                Sequence 45, Application US/09464453 Patent No. 6358686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 02-DEC-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 07-MAR-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 415 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Brassica Polymorphisms NUMBER OF SEQUENCES: 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: replace(21, "")
OTHER INFORMATION: /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 415 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Liebeschuetz, Joe REGISTRATION NUMBER: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
                                                                             ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                               Landry, Benoit S.
Sapolsky, Ronald J.
TITLE OF INVENTION: Brassica Polymorphisms
                                                                                                                                                                                                                                                                                                                                       APPLICANT: Lemieux, Bertrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    32 TTGCGACTCTGTGGGACTGTAACTGTC 6
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                 SOFTWARE: FastSEQ for Windows Version 2.0
                                        OPERATING SYSTEM: DOS
                                                                COMPUTER:
                                                                                                                                                             STATE: California
                                                                                                                                                                                CITY: San Francisco
                                                                                                                                                                                                 STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      California
                                                                                                                                                                                                                       ADDRESSEE: Townsend and Townsend and Crew LLP
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02-DEC-1996
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                                                       IBM Compatible
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APPLICANT: Harkki, Anu M.
                                                   APPLICATION NUMBER: US 08/
FILLING DATE: 24-AUG-1993
APPLICATION NUMBER: US 07/
FILLING DATE: 05-NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 45:
                                                                                                                                                                                                                                                                                                                                   ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Harkki, Anu M.
APPLICANT: Myasnikov, Andrey N.
APPLICANT: Apajalahti, Juha H.A.
APPLICANT: Bastinen, Ossi A.
TITLE OF INVENTION: Manufacture of Xylitol
                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Pelacity
                                                                                                                                                                                                                       FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                      REGISTRATION NUMBER: 33,851
                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Washington
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OTHER INFORMATION: /note= "deletion polymorphism"
SEQUENCE DESCRIPTION: SEQ ID NO: 45:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: 415 576-0200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   32 TTGCGACTCTGTGGGACTGTAACTGTC 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      43 TTTCAACACTATCGGAATGTGACTGTC 69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/464,453 FILING DATE: 14-Dec-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 018547-030100US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     relex: <Unknown>
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 Mismatches

                   1427.0010001
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                                                                        Matches
                                                                                                        Query Match
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Best Local Similarity 74.1%;
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TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                  TELEFAX: 310-445-9031
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                           TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 310-445-1140
                                                                                                                                                                                                                                                                                                                                                                        NAME: Adriano, Sarah B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: No. 5728821el Mutant BR96 Antibodies and TITLE OF INVENTION: Functional Equivalents Reactive With Human Carcinomas NUMBER OF SEQUENCES: 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
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                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 304
LECOMMUNICATION THRONG
                                                                                                                                                                                                                                                                                                                                                  NAME: Adriano, Sarah B
REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/0
FILING DATE: 04-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STATE: CA
                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
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STRANDEDNESS: both
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                                  70 GGGCCTCCAGAAAACATTGAGG 91
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2 GGGACTCCAGAAAGCTTTTAGG 23
                                                                    18; Conservative
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                                                                                     h 16.8%;
Similarity 81.8%;
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Huse, William
Rosok, Mae J.
                                                                                                                                                                                                                              33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Merchant & Gould
11150 Santa Monica Bldv., Suite 400
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                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PatentIn Release #1.0, Version #1.25
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                                                                                                                                                         DNA (genomic)
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                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                      30436.16US01
                                                                                 Score 15.6; DB 1; Length 33; Pred. No. 1.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.8; DB 1;
Pred. No. 1.6e+03;
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                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
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RESULT 18

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                                                                                                                                                                                                                                                                                                                                                                               ; Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-487-860-14
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; Patent No. 5792456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-487-860-14
                                                                                                                                                                                                                                                                                                                                                            ; GENERAL INFORMATION:
                   EARLIER APPLICATION NUMBER: 60/069,865
EARLIER FILING DATE: 1997-12-17
EARLIER APPLICATION NUMBER: 08/866,340
EARLIER FILING DATE: 1997-05-30
NUMBER OF SEQ ID NOS: 138
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                           TITLE OF INVENTION: DNA METHYLTRANSFERASE GENOMIC SEQUENCES AND ANTISENSE TITLE OF INVENTION: OLIGONUCLEOTIDES
                                                                                                                                                                                                                                                                                      APPLICANT: Szyf, Moshe
APPLICANT: Bigey, Pascal
APPLICANT: Ramchandani, Shyam
                                                                                                                                                                          FILE REFERENCE: 106101.194
CURRENT APPLICATION NUMBER: US/09/103,875A
CURRENT FILING DATE: 1998-06-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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LENGTH: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 34,470
REFERENCE/DOCKET NUMBER: 30.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 310-445-1140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Huse, William
APPLICANT: Rosok, Mae J.
TITLE OF INVENTION: No. 5792456el Mutant BR96 Antibodies and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Yelton, Dale APPLICANT: Glaser, Scot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: Adriano, Sarah B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: No. 5792456el Mutant BR96 Antibodies and TITLE OF INVENTION: Functional Equivalents Reactive With Human Carcinomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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CA
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                                                                                                                                                                                                                                                                                                                                                                                                     Application US/09103875A
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81.8%;
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            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: LARRY GOLD et al.
TITLE OF INVENTION: HIGH AFFINITY OLIGONUCLEOTIDE
TITLE OF INVENTION: LIGANDS TO GROWTH
                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                              FILING DATE: 20-MARCH-19: ATTORNEY/AGENT INFORMATION:
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CITY: Englewood
CITY: Colorado
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                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 02-JUN
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                                                                                                                 REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/479,725
                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/973,124
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                                                                TELEPHONE:
                                                                                                 REFERENCE/DOCKET NUMBER
                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/479,783
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21;
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Similarity 70.0%;
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                                                                                                                               Barry J. Swanson
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41 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8400 E. Prentice Avenue, Suite 200
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                                                  (303)
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WordPerfect 6.1
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                                                                (303) 793-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JUNE-1995
                                                                                                                                                                                                               07-JUNE-1995
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                                                                                                                                                                  20-MARCH-1996
                                                                                                                                                                                                                                                                 07-JUNE-1995
                                                  793-3433
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                                                                                                                                                                                                                                                                                                                                    08/465,591
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                                                                                                                                                                                08/618,693
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                                                                                                                 33,215
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US-08-973-124-285
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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                                                                                                                                                              FILING DATE: 07-JUNE-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                 FILING DATE: 07-JUNE PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                         FILING DATE: 05-JUNE-1995
PRIOR APPLICATION DATA:
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                                                                                                       FILING DATE: 20-MARCH-19 ATTORNEY/AGENT INFORMATION:
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             TELEPHONE: (303) 793-3333
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                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
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SOFTWARE: WordPerfect 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
                                                                    NAME: Barry J. Swanson REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                        FILING DATE:
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                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                APPLICATION NUMBER:
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                                                                                                                                           APPLICATION NUMBER:
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Colorado
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(303) 793-3433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LARRY GOLD; NEBOJSA JANJIC; STEVEN RINGQUIST; PAGRATIS; PENELOPE J. TOOTHMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Swanson & Bratschun, L.L.C
                                                                                                                                                                                                                                                                                                                                                                                                   02-JUNE-1995
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                                                                                                                                                                                                                                                                                        05-JUNE-1995
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                                                                                                                                                                                                                                   07-JUNE-1995
                                                                                                                           20-MARCH-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIGH AFFINITY OLIGONUCLEOTIDE LIGANDS TO TRANSFORMING GROWTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FACTOR (TGF ), PLATELET-DERIVED GROWTH FACTOR (PDGF) AND HUMAN KEROTINOCYTE GROWTH FACTOR (hKGF)
                                                                                                                                                                                                08/479,783
                                                                                                                                                                                                                                                    08/479,725
                                                                                                                                                                                                                                                                                                         08/465,591
                                                                                                                                                                                                                                                                                                                                                              08/465,594
                                                                                                                                                                                                                                                                                                                                                                                                                     08/458,424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08/458,423
                                                                                                                                             08/618,693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 4; Length 41; Pred. No. 2.3e+03;
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US-08-340-426D-63/c
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                                     RESULT 23
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US-08-454-557C-63/c
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Sequence 63, Application US/08340426D
                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack R.
TITLE OF INVENTION: of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                        TYPE: nucleic acid STRANDEDNESS: both TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0609.3840003
                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 30-MA)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity es 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           THER INFORMATION: All pyrimidines are 2'-fluoro (2'-F) modified
                                                                                       27 TTCAGCCTGGGTGACAGAGC 8
                                                                                                                           18 TTCTGCCTGGGTGACAATGC 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30 GACAATGCCTTGGTTTCAACACTAT 54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          D.C
                                                                                                                                                                                                                                                                                                      31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Sterne, Kessler, Goldstein & Fox P.L.L.C
1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                               (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                Conservative
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                                                                                                                                                                             16.3%; Score 15.2; DB 2; Length 31; 85.0%; Pred. No. 2.5e+03;
                                                                                                                                                                                                                                                                                                                                                                               371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.6%; Score 15.4; DB 5; 52.0%; Pred. No. 2.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US/08/454,557C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
                                                                                                                                                                Mismatches
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                                                                                                                                                              Indels
                                                                                                                                                            0;
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                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 63,
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                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: de la Monte, SI
APPLICANT: Wands, Jack R.
                                                        CURRENT APPLICATION DATA
                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection TITLE OF INVENTION: of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection TITLE OF INVENTION: of Alzheimer's Disease NUMBER OF SEQUENCES: 121
                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
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CLASSIFICATION: 530
                                                                     COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                  APPLICATION NUMBER: US/08/450,673C FILING DATE: 30-MAY-1995
                                                                                                                                                                              ZIP: 20005-3934
                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                 CITY: Washington
                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: both
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REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                  D.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08450673C
                                                                                                                                                                                                                                                      E: Sterne, Kessler, Goldstein & Fox P.L.L.C. 1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        : (202) 371-2600
(202) 371-2540
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                                                                                                                                                                                                                                                                                                                                                                                          Suzanne
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; Sequence 63, Applicat
; GENERAL INFORMATION:
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PCT-US95-17111A-63/c
                                                                                                                              PCT-US95-17111A-63
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                                                          Matches
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TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 63
                                                                                                                                                                                                                                       TELEFAX: (202) 371-25-
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 371-2600 TELEFAX: (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/340,426
FILING DATE: 14-NOV-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Neural Thread Protein Gene Expression and TITLE OF INVENTION: Detection of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: de la Monte, Suzanne APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1100
CTTY: Washington
                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                          TYPE:
                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                     NAME: Ludwig, Steven R. REGISTRATION NUMBER: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US95/17111A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Ludwig, Steven R. REGISTRATION NUMBER: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                          Local Similarity
                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                    LENGTH:
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                      18 TTCTGCCTGGGTGACAATGC 37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17; Conservative
                                                        17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RY: U.S.A.
20005-3934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 16.3%;
Similarity 85.0%;
                                                                                                                                                                                    nucleic acid
Application PC/TUS9517111A
                                                                                                                                                                                                    31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31 base pairs
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                                                                                                                                                  hoth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sterne, Kessler, Goldstein & Fox P.L.L.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                    both
                                                                        16.3%; Score 15.2; DB 5; Length 31; 85.0%; Pred. No. 2.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121
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                                                      0; Mismatches

 Mismatches

                                                                                                                                                                                                                                                                                                                   0609.3840002
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                                                        3; Indels
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                                                    0;
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LENGTH: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application PC/TUS9414888 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/344,700 CURRENT FILING DATE: 1999-06-25 EARLIER APPLICATION NUMBER: PCT/JP97/0485 EARLIER FILING DATE: 1997-12-25 EARLIER APPLICATION NUMBER: JP 8/357864 EARLIER FILING DATE: 1996-12-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: NOVEL SERINE-THREONINE KINASE GENEFILE REFERENCE: 06501/033001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Nezu, Jun-
APPLICANT: Oku, Asuka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 37
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                        INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                          REFERENCE/DOCKET NUMBER: FP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Homologous Recombination for Universal TITLE OF INVENTION: Donor Cells and Chimeric Mammalian Hosts
                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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MOLECULE TYPE:
                                                                                        SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE:
                              STRANDEDNESS:
                                                                    LENGTH:
                                                                                                                              TELEFAX:
                                                                                                                                                                                                  REGISTRATION NUMBER:
                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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                  TOPOLOGY:
                                                                                                                                                                                                                      NAME:
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Local Similarity 63.9%;
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                                                  nucleic acid
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                                                                                                                                                                                                                      Rowland,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09344700
                                                                    50 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                            415-398-3249
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                                                                                                                                                                                                                                                                                                                                                                 IBM PC compatible
                  linear
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Pred. No. 2.7e+03;
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US-07-771-022F-8/c
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Patent No. 5476926
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                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617) 632-4012
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: DFCI-117
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 632-4016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Adipocyte-Specific DNA
TITLE OF INVENTION: Sequences And Use Thereof In The Production Of
TITLE OF INVENTION: Transgenic Animals Exhibiting Altered Fat
TITLE OF INVENTION: Tissue Metabolism.
                                                                                           IMMEDIATE SOURCE:
LIBRARY: genom
                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07/589,971
FILING DATE: 28-SEP-1990
ATTORNEY/AGENT INFORMATION:
NAME: HART, JULIA D.
REGISTRATION NUMBER: 33132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: WordPerfect 6.0
CURRENT APPLICATION DATA:
                                                        FEATURE
                                                                                                                                                                                                                                               ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                 HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                  TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: GRAVES, REED A APPLICANT: ROSS, SUSAN R.
                                                                                                                                                                                                                                                                   ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                  NAME/KEY:
                                                                  CLONE: aP2911
                                                                                                                             CELL TYPE: fibroblast CELL LINE: 3T3-F442A
                                                                                                                                                                                      STRAIN: Swiss Webster DEVELOPMENTAL STAGE: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette, 3.50 inch, 1 MB storage COMPUTER: IBM-compatible Compaq Prolinea MT 4/66 OPERATING SYSTEM: MS/DOS COMPUTER: WGYADAFFOCK 6
LOCATION: 8 to 19
                                                                                                                                                                      TISSUE TYPE:
                                                                                                                                                                                                                             ORGANISM:
                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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Similarity 71.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPIEGELMAN, BRUCE M.
                                  aP2 regulatory element; identified herein as
                                                                                                                                                                                                                             Mus musculus
                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                    fibroblast
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                                                                                                                                                                      embryonic fibroblast
                                                                                                                                                                                                                                                                                                    genomic DNA
                                                                                                                                                                                                                                                                                                                                          double
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US-07-771-022F-8
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; Patent No. 5981262
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Brugge, Joan APPLICANT: Morgenstern, APPLICANT: Shiue, Lily
                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO US94/04540
                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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TELECOMMUNICATION INFORMATION:
                                                      ATTORNEY/AGENT INFORMATION:
NAME: BERSTEIN, David L.
                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Pawson, Anthony TITLE OF INVENTION: HUMAN:
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                                                                                      CLASSIFICATION: 435
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DATE: 00-MAR-1992
                                  REGISTRATION NUMBER:
                                                                                                          APPLICATION NUMBER: FILING DATE: 23-API
                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 31-JA
                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Cambridge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE: Analysis of a Tissue-Specific Enhancer: ARF6 TITLE: Regulated Addipogenic Gene Expression
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IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
IDENTIFICATION METHOD:
                  REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AUTHORS:
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                                                  BERSTEIN, David
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26 Landsdowne Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Morgenstern, Jay
Shiue, Lily
                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zydowsky, Ly
Zoller, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SPEIGELMAN, BRUCE M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TONTONOZ, PETER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GRAVES, REED A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecular and Cellular Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ARIAD Pharmaceuticals, Inc.
                                                                                                                                                                                     25-APR-1994
                                                                                                                                                                                                                                                           31-JAN-1996
                                                                                                          23-APR-1993
                                                                                                                                                                                                                                                                                                                                                                          Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HUMAN syk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lynne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        correspond to nucleotides 93 to 104 of the 518 bp enhancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the nuclear extract of differentiated adipocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 1; this sequence binds a protein factor present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleotides 8 to 19 of SEQ ID NO 6
                                                                                                                              us 08/052,560
                                                                                                                                                                                                                                                                               us/08/596,319
                              31,235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in pSVKS1-CAT vector; transfection assays; gel mobilit shift
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                  ARIAD305A-PCT/US
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 12, Application US/07771022F Patent No. 5476926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                   TELEPHONE: (617) 632-4016
TELEFAX: (617) 632-4012
TELEFAX: (617) 630-4012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
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TELEFAX: (617) 494-0208
INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ROSS, SU
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WordPerfect 6.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
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                                                                 ORIGINAL SOURCE:
                                                                                           HYPOTHETICAL:
                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
          STRAIN: Swiss Webster DEVELOPMENTAL STAGE: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette, 3.50 inch, 1 MB storage COMPUTER: IBM-compatible Compaq Prolinea MT 4/66 OPERATING SYSTEM: MS/DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
TISSUE TYPE: embryonic fibroblast
                                                ORGANISM:
                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/07/771,022F
FILING DATE: 07-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/589,971 FILING DATE: 28-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                          LENGTH:
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                                                                               SENSE:
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                                                                                                                                                                                                                                                                                                     HART,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32 base pairs
                                                                                                                                                                        33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       44 Binney Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VENTION: Adipocyte-Specific DNA
VENTION: Roll Production of VENTION: Sequences And Use Thereof In The Production of VENTION: Transgenic Animals Exhibiting Altered Fat VENTION: Tissue Metabolism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SPIEGELMAN, BRUCE M.
                                              Mus musculus
                                                                                                                             linear
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                                                                                                                                                                                                                                                                             JULIA D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The Dana-Farber Cancer Institute
                                                                                              no
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                                                                                                              genomic DNA
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                                                                                                                                           double
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75.0%;
               embryonic
                                                                                                                                                                                                                                                                     DFCI-117
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Pred. No. 3e+03;
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RESULT 32
US-09-237-712-10
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                                                            ; Sequence 10, Application US/09237712
; Patent No. 6180391
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US-09-237-712-9/c
                                                                                                                                                                                                                                                                                                                                   US-09-237-712-9
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                                         ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                           Query Match
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APPLICANT: BROWN, WILLIAM C.
TITLE OF INVENTION: HIGHLY EFFICIENT CONTROLLED EXPRESSION OF EXOGENOUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/237,712 CURRENT FILING DATE: 1999-01-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: A-518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: HIGHLY EFFICIENT CONTROLLED EXPRESSION OF EXOGENOUS TITLE OF INVENTION: GENES IN E. COLI
                                                                                                                                                                                                                                                                                                                                                                                                                                            EARLIER APPLICATION NUMBER: 60/072,794
EARLIER FILING DATE: 1998-01-28
NUMBER OF SEQ ID NOS: 98
                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                 ORGANISM: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PUBLICATION INFORMATION:
AUTHORS: GRAVES, REED A.
                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AUTHORS:
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OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CELL TYPE:
CELL LINE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE: Analysis of a Tissue-Specific Enhancer:
                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JOURNAL:
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                                                                                                                                                                             35 GAGTTGGAAGAAATCCATTCAGC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        33 CAATCCCTTCTCTCTGGGTGAAA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11 CACTCCATTCTGCCTGGGTGACA 33
                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                 1 GAGCTGGGAGCACTCCATTCTGC 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BROWN, WILLIAM C.
                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TONTONOZ, PETER
SPEIGELMAN, BRU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3T3-F442A
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                                                                                                                                                                                                                                                                      16.1%;
78.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleotide sequence of SEQ ID NO 8 herein and includes an additional nine nucleotides, "TCGGCGCAC" of 5' flanking region of the mouse aP2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the aP2 regulatory element identified herein as ARE6. The sequence encompas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This nucleotide sequence includes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BRUCE M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                         Score 15; DB 4;
Pred. No. 3.2e+03,
                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                         DB 4; Length 36;
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US-08-191-866D-91
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                                                        Matches
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CURRENT FILING DATE: 1999-01-26
EARLIER APPLICATION NUMBER: 60/072,794
EARLIER FILING DATE: 1998-01-28
NUMBER OF SEQ ID NOS: 98
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
                                                                                       Query Match
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                                                                                                                                                                                                                                                            TELEX: 422523
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Cochran, Mark D
APPLICANT: Macdonald, Richard D.
                                                                                                                                                         HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                               TELEPHONE: (212) 278-0400
                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Recombinant Infectious Bovine TITLE OF INVENTION: Rhinotracheitis Virus S-IBR-052 And Uses Thereof
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: New York
                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/191,866D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                        Local
                                                                                                                                                                                                                                                                                                                  TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             POCGI
                                                                                                                                                                                                                                              ENGTH:
                33 AATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GAGCTGGGAGCACTCCATTCTGC 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 GAGTTGGAAGAAATCCATTCAGC 30
                                                  21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         h 16.1%; Score 15; DB 4;
Similarity 78.3%; Pred. No. 3.2e+0;
18; Conservative 0; Mismatches
                                                                     Similarity
                                                                                                                                                                                                                            nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                   White, John P.
New York
                                                                                                                                                                                                                                         41 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                              (212) 391-0525
                                                                                                                                          NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              John P. White
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                                                                                                                                                                                                                                                                                                                                                                                                                                  4 February 1994
                                                                                                                                                                         DNA (genomic)
                                                                                                                                                                                                             double
                                                                 16.1%;
67.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99
                                                                                                                                                                                                                                                                                                                                                                    28,678
                                                                                                                                                                                                                                                                                  91:
                                                    0; Mismatches
                                                                   Score 15; DB 1;
Pred. No. 3.4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Version #1.25
                                                                                    DB 1; Length 41;
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                                                    10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 36;
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                                                                                                                                                                                                                                                                          RESULT 35
                                                                                                                                                                                           ; Sequence 143, Application US/08295802; Patent No. 6127163; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-480-640A-143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 143, Application US/08480640A Patent No. 6033904
                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: White, John P
REGISTRATION NUMBER: 28,678
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Cochran, Mark D. APPLICANT: Junker, David E. TITLE OF INVENTION: Recombin NUMBER OF SEQUENCES: 225
                                                                                                          CORRESPONDENCE ADDRESS
                                                                                                                        TITLE OF INVENTION: Recombinant Swinepox Virus NUMBER OF SEQUENCES: 188
                                                                                                                                                             APPLICANT: Cochran Ph.D., Mark I APPLICANT: Junker M.S., David E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: I
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ATTORNEY/AGENT INFORMATION:
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                                                   STREET:
ZIP: 10112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Bovine respiratory syncytial virus STRAIN: Strain 375
                   COUNTRY:
                                 STATE: New York
                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 07-JU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                    Match 16.1%; Score 15; DB 3; Local Similarity 67.7%; Pred. No. 3.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH: 41 base pairs
                                                                                                                                                                                                                                                                                                                                 40 AATGTCTTGAATTTAAGATGATGGGTATGAG 10
                                                                                                                                                                                                                                                                                                                                                                   33 AATGCCTTGGTTTCAACACTATCGGAATGTG 63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid
                                                   New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: John P. White
1185 Avenue of the Americas
                                                                     30 Rockefeller Plaza
                 USA
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                                                                                      John P. White
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
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                                                                                                                                                                                   Mark D
                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 41;
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             ; Sequence 143, Application US/08488237A
; Patent No. 6251403
                                                     US-08-488-237A-143/c
                                                                       RESULT 37
                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: primer US-08-686-968C-70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; GENERAL INFORMATION:
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                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/08/686,968C CURRENT FILING DATE: 1996-07-25 NUMBER OF SEQ ID NOS: 231
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Junker, David E.
TITLE OF INVENTION: Recombinant Swinepox Virus
FILE REFERENCE: 39119-H/JML
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cochran, Mark D. APPLICANT: Junker, David I
                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 41
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NAME: White, John P
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: double
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OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEFAX: (212)664-0525
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 16.1%; Score 15; DB 3; Local Similarity 67.7%; Pred. No. 3.4e+03;
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                                                                                                                                                                                                                      Local
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                                                                                                                         40 AATGTCTTGAATTTAAGATGATGGGTATGAG 10
                                                                                                                                                             33 AATGCCTTGGTTTCAACACTATCGGAATGTG 63
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                                                                                                                                                                                                                  h 16.1%; score 15; DB 4; similarity 67.7%; Pred. No. 3.4e+03
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                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bovine respiratory syncytial virus
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 Mismatches

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                                                                                                                                                                                                 10; Indels
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                                                                                                                                                                                                                                                                                                                                                                                           Sequence 30, Application US/08672213 Patent No. 6306649
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TELEPHONE: (212) 278-0400
TELEPAX: (212) 391-0525
INFORMATION FOR SEQ ID NO: 143:
            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                            ZIP: 02139-4234
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSE: ARIAD Gene Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: USE OF HETEROLOGOUS TRANSCRIPTION TITLE OF INVENTION: FACTORS IN GENE THERAPY
                                                                                                                                                                                                                                                                                                                              APPLICANT: GILMAN, Michael Z. APPLICANT: NATESAN, Sridaran
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ANTI-SENSE: 1
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APPLICANT: Junker, David E.
TITLE OF INVENITION: Recombinant Swinepox Virus
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: Patentl
  APPLICATION NUMBER:
                                                                                                                                                                                         CITY: Cambridge
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                       COUNTRY:
                                                                                                                                                                               STATE:
                                                                                                                                                                                                                STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/488,237A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         40 AATGTCTTGAATTTAAGATGATGGGTATGAG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               33 AATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      White, John P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
                                                                                                                                                                        Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l: Bovine respiratory syncytial virus
Strain 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41 base pairs
                                                                                                                                                                                                              26 Landsdowne Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1185 Avenue of the Americas
                                                                                                                                                       USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16.1%; Score 15; DB 4; Length 41; 67.7%; Pred. No. 3.4e+03;
US/08/672,213
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    Mismatches

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RESULT 39
US-08-375-992A-143/c
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US-08-672-213-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 143, Application US/08375992A Patent No. 6328975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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INFORMATION FOR SEQ ID NO: 30:
                                                                 TELEFAX: (212) 391-0525
INFORMATION FOR SEQ ID NO: 143:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cochran, Mark D.
APPLICANT: Junker, David E.
TITLE OF INVENTION: Recombinant Swinepox Virus
NUMBER OF SEQUENCES: 220
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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LENGTH: 41 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 60 FILING DATE: 29-DEC-1995 ATTORNEY/AGENT INFORMATION:
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                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 278-0400
                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
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                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      STATE: New York
                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
COMPUTER: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
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                 STRANDEDNESS:
                                                                                                                                                            NAME: White, John P
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY:
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REFERENCE/DOCKET NUMBER: ARIAD 346B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
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TOPOLOGY:
                                TYPE: nucleic acid
                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
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                                                    LENGTH: 41 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                               New York
                                                                                                                                                                                                                                                                                                                                                                                                                                              1185 Avenue of the Americas
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linear
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                                                                                                                                                                                                                                 Herewith
                                                                                                                                                                                                                                                                                                                                        Floppy disk
                 double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16.1%; Score 15; DB 4; 67.7%; Pred. No. 3.4e+03
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                                                                                                                                                              28,678
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יייטאייזאא: Bovine respiratory syncytial virus
; STRAIN: Strain 375
US-08-375-992A-143
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                                                               US-08-596-387B-38
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 Matches
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               Best Local Similarity
                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                   TELEFAX: (708) 267-5376 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                      FILING DATE: 01-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/283,302
FILING DATE: 29-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: DESCRIPTION
                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/I
FILING DATE: 31-JUL-1995
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APPLICANT: Jiao, J
                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                               NAME: Pearson, Louise S. REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                   TYPE:
                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                   NAME:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
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                                                                                                                                 LENGTH:
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                                                                                 POPOLOGY:
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21; Conservative
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                                                                                                              : 42 base pairs
nucleic acid
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1717 Deerfield Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
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Burkhardt, Martin
                                                                                 unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PatentIn
                                                                                                                                                                                               (708) 267-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NO
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                                                                                               unknown
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67.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peter R.
                 16.1%; Score 15; DB 2; 67.7%; Pred. No. 3.4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                 PCT/US95/09816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                               US 08/382,454
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 0; Mismatches
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Pred. No. 3.4e+0;
                                 Length 42;
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 0
 Gaps
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11 CACTCCATTCTGCCTGGGTGACAATGCCTTG 41

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1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31

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PCT-US95-09816A-38
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                                                                                    Sequence 38, Application PC/TUS9509816A GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                      Query Match
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Patent No. 6309645
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                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                  APPLICANT:
APPLICANT:
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (708) 267-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 29-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 01 FILING DATE: 01-FEB-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Burkhardt, Martin APPLICANT: Wong, Hing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Rhode, Peter R. APPLICANT: Jiao, Jin-An
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: MHC COMPLEXES AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: unkno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Pearson, Louise S.
REGISTRATION NUMBER: 32,369
REFERENCE/DOCKET NUMBER: ST
                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/596,387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/067,615
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                11 CACTCCATTCTGCCTGGGTGACAATGCCTTG 41
                                                                                                                                                                                   1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31
                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09067615
                                                                Wong, Hing C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1717 Deerfield Road
Edwards, Ana C
                                                Rhode, Peter R.
                                                                                                                                                                                                                                                                                                                                                                                                                                      (708)
                Grammer, Susan
                                   Widanz, Jon A.
                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                     unknown
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                                                                                                                                                                                                                                                                                                                                                  unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                      267-5376
                                                                                                                                                                                                                                                                   16.1%; Score 15; DB 4; 67.7%; Pred. No. 3.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US 08/283,302
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                                                                                                                                                                                                                                                                                                                                                                                                                        38:

 Mismatches

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RESULT 43
US-08-681-935-13
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                                                                                                                                                                                                                                                                                                                                                           Patent No. 5710248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                              Sequence 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                             GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (708) 267-5376 INFORMATION FOR SEQ ID NO: :
          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
COMPATING SYSTEM: PC-DOS/MS-DOS
                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                               APPLICANT: GROSE, CHARLES F.
TITLE OF INVENTION: PEPTIDE TAG FOR IMMUNODETECTION AND
TITLE OF INVENTION: IMMUNOPURIFICATION
NUMBER OF SEQUENCES: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: PC'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Chavaillaz, Pierre-Andre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICATION NUMBER:
                                                                                                                                                 COUNTRY:
                                                                                                                                                                       STATE:
                                                                                                                                                                                                        STREET:
                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity les 21; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Pearson, Louise S. REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 01-FEB-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11 CACTCCATTCTGCCTGGGTGACAATGCCTTG 41
                                                                                                                                                                                                                                                                                                                                         INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CCCCCCAGGCTTCCCGGGCCACCATGCCGTG 31
                                                                                                                                                                                        MINNEAPOLIS
                                                                                                                                                                   MINNESOTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Illinois
                                                                                                                                                                                                                                                                                                                                                                              Application US/08681935
                                                                                                                                                                                                        E: MUETING, RAASCH, GEBHARDT & SCHWAPPACH, P.A. 119 NORTH FOURTH STREET, SUITE 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Dade International, Inc.
1717 Deerfield Road
                                                                                                                                                   USA
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                                                                                       Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unknown
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US/08/681,935
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US-08-939-323-13
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                                                       US-08-939-323-13
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Best Local Similarity
                 Query Match
                                                                                                                                                                               NAME: MUETING MS., ANN M.
REGISTRATION NUMBER: 33,977
REFERENCE/DOCKET NUMBER: 140.00040102
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 305-1217
TELEPHONE: (612) 305-1228
INFORMATION FOR SEQ ID NO: 13:
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                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 45 base pairs
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/681,935 FILING DATE: 29-JUL-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: GROSE, CHARLES F.
TITLE OF INVENTION: PEPTIDE TAG FOR IMMUNODETECTION AND TITLE OF INVENTION: IMMUNOPURIFICATION
NUMBER OF SEQUENCES: 21
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                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
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ADDRESSEE: MUETING, RAASCH & GEBHARDT, P.A
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                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                         TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30 GACAATGCCTTGGTTTCAACACTATCGGAATGTGACTGT 68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              119 NORTH FOURTH STREET, SUITE 203
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                                                                       other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                        29-SEP-1997
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16.1%;
61.5%;
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Score 15; DB 4;
Pred. No. 3.6e+03;
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                 Length 45;
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RESULT 46
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                                                                                                                            Matches
                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: TREATMENT OF DISEASES OR TITLE OF INVENTION: CONDITIONS RELATED TO LEVEL TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/005,974 FILING DATE: October 26, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 27 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                       FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Los Angeles
STATE: California
                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warburg, Richard REGISTRATION NUMBER: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: JOCAL CLASSIFICATION:
                                                                                                                                                                                                                 OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30 GACAATGCCTTGGTTTCAACACTATCGGAATGTGACTGT 68
                                                                                        61 GTGACTGTCGGGCCTCCAGAAAACATT 87
                                                     27 GTGAGTGTTTCNTCATCAGAAAACATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7 GTCACCGTATTGCCTTTGACAGTATGCGATTGTGATAGT 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            McSwiggen, James
Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Escobedo,
                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/08584040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     January 11, 1996
                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GROWTH FACTOR
                                                                                                                                          15.9%; Score 14.8; DB 4; Length 27; 70.4%; Pred. No. 3.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM P.C. DOS 5.0
                                                                                                                                                                                                                 The letter "N" represents the stem II region of an HH ribozyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US/08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15; Indels
                                                                                                                              8
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US-08-584-040-5068/c

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                                                                                                                                                                    RESULT 47
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                                                                                                           Sequence 33, App. 5-06 No. 598559
                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5068, Application US/08584040 Patent No. 6346398
                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 27 base pairs
                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/005,97. FILING DATE: October 26, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 514 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
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 APPLICANT:
                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: The letter "N" represents the stem II region OTHER INFORMATION: of an HH ribozyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/584,040 FILING DATE: January 11, 1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
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                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                    61 GTGACTGTCGGGCCTCCAGAAAACATT 87
                                                                                                                                                                                                                     27 GTGAGTGTTTCNTCATCAGAAAACATT 1
                                                                                                                                                                                                                                                                                       19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     California
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                                                                                                                                 Application US/08332562A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Escobedo,
                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pavco, Pamela
HULETT, Mark D.
                               HIBBS, Margaret L. SCOTT, Bernadette M.
                                                               MCKENZIE, Ian F.C. HOGARTH, Mark P.
                BONADONNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                                                                                                    15.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      METHOD AND REAGENT FOR THE
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    Mismatches

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                                                                                                                                                                                                                                                                                                    Score 14.8; DB 4;
Pred. No. 3.4e+03;
                                                                                                                                                                                                                                                                                       8; Indels
                                                                                                                                                                                                                                                                                                                    Length 27;
                                                                                                                                                                                                                                                                                   0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 147, Application US/08813507; Patent No. 6114116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
                                                                                                                                                          ZIP: 94111-3834
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                        APPLICANT: Sapolsky, Ronald J. TITLE OF INVENTION: Brassica Polymorphisms NUMBER OF SEQUENCES: 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07/896,457 FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/0 FILING DATE: 31-OCT-1994
                                                                                   SOFTWARE:
                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                               FILING DATE: 07-MAR-1997
                                                                                                   OPERATING SYSTEM:
                                                                                                                         COMPUTER:
                                                                                                                                                                                                                 STATE:
                                                                                                                                                                                                                                                                   ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
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                 CLASSIFICATION:
                                                                                                                                          MEDIUM TYPE:
                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 15.9%;
Local Similarity 73.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             44 TTCAACACTATCGGAATGTGACTGTC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 TTCTAGACTATTGGACAGTGATGGTC 28
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                                                                                                                                                                                                                                San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  D.C
                                                                                                                                                                                                                 California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Foley & Lardner
3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                    Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                       Lemieux, Bertrand
Landry, Benoit S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (202)672-5399
                                                                                     FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                     IBM Compatible
                                                                                                                                                                                                                                                                     Townsend and Townsend and Crew LLP
                                                                                                                                          Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US/08/332,562A
                                                  US/08/813,507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8;
Pred. No. 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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; SEQUENCE DESCRIPTION: SEQ ID NO: 147: US-09-464-453-147
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US-09-464-453-147
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 147,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 147:
                                                                                                                           INFORMATION FOR SEQ ID NO: 147: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 41 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 415 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Liebeschuetz, Joe
REGISTRATION NUMBER: 37,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 15.9%; Score 14.8; DB 3; Length 41; Local Similarity 61.1%; Pred. No. 4.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 018547-030100US TELECOMMUNICATION INFORMATION: TELEPHONE: 415 576-0200
                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/813,507
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Brassica Polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lemieux, Bertrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28 GTGACAATGCCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 GTGAAAAGGCATTGTKCTCCAAAATCTCGCTCTGCG 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22; Conservative
                                                           LENGTH: 41 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                            NAME: Liebeschuetz, Joe REGISTRATION NUMBER: 37
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/464,453
FILING DATE: 14-Dec-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                        TELEX: <Unknown>
                                                                                                                                                                                          TELEFAX: 415 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09464453
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Landry, Benoit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US 60/032,069

    Mismatches

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US-07-963-331D-5
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; Patent No. 5334511
                                                                         Matches
                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Harada, Yasuhiro APPLICANT: Nakano, Elichi APPLICANT: Tatsumi, Hiroki APPLICANT: Umezu, Motoaki TITLE OF INVENTION: Hormone (TITLE OF INVENTION: Methods! NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 433-87
TELEX: 278356
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 31,243
REFERENCE/DECKET NUMBER: HI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 433-4150
TELEFAX: (415) 433-8716
                                                                                                                                                                                                                                                      TOPOLOGY: Linear MOLECULE TYPE: Othe
                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                      MOLECULE TYPE: DNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                              OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 2002
CITY: San Francisco
                                                                                         Match 15.9%; Local Similarity 73.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 19921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 15.9%;
Local Similarity 61.1%;
                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
19 TCCAGCCTGTTTGCCAATGCCGTGCT 44
                     18 TTCTGCCTGGGTGACAATGCCTTGGT 43
                                                                         19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28 GTGACAATGCCTTGGTTTCAACACTATCGGAATGTG 63
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                                                                                                                                                                                                                                                                                                        NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dergosits, Michael E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001 Ferry Building
                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Limbach & Limbach
                                                                                                                                                                                                                                                      Other nucleic acid (synthesized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19921016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                                                                                                                                                                                                                       Single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated Mink Growth
Hormone Genes, No. 5334511el Recombinant DNA And
Methods For Producing Mink Growth Hormone.
                                                                                                                                                                hormone
                                                                                                                                                                                Based on the base sequence of rat growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US/07/963,331D
                                                                         0; Mismatches

    Mismatches

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                                                                                           Score 14.8; DB 1;
Pred. No. 4.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 4; Length 41; Pred. No. 4.1e+03;
                                                                                                          Length 45;
                                                                         Indels
                                                                         0,
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                                                                         Gaps
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US-09-167-681-6

RESULT 51

; Sequence 6, Application US/09167681A

Patent No.

6265561

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; Sequence 23, Application US/08557128
; Patent No. 5849524
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 62655
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Weinshilboum, M.D., Richard M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Weinshilboum, M.D., Richard M.
APPLICANT: Raftogianis, Rebecca B.
APPLICANT: Wood, Thomas C.
APPLICANT: Ofterness, Diane M.
TITLE OF INVENTION: SULFOTRANSFERASE SEQUENCE VARIANTS
FILE REFERENCE: 07039/118001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: KONDO, Keiji
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: WOOD, Thomas C.
APPLICANT: Otterness, Diane M.
TITLE OF INVENTION: SULFOTRANSFERASE SEQUENCE VARIANTS
FILE REFERENCE: 07039/118001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/167,681A
CURRENT FILING DATE: 1998-10-07
NUMBER OF SEQ ID NOS: 52
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CURRENT FILING DATE: 1998-10-07
NUMBER OF SEQ ID NOS: 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Raftogianis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 3.0
APPLICANT: KONDO, Keiji
APPLICANT: KAJIWARA, Susumu
APPLICANT: MISAWA, No. 58495241hiko
APPLICANT: MISAWA, NO. 58495241hiko
TITLE OF INVENTION: TRANSFORMATION SYSTEMS FOR THE YEAST
TITLE OF INVENTION: CANDIDA UTILIS AND THE EXPRESSION OF HETEROLOGOUS GENES
TITLE OF INVENTION: THEREWITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 25
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                  15 CCATTCTGCCTGGGTGACAAT 35
                                                                                                                                                                                                                                                                                                                                        1 CCACTCTGCCTGGCCCACAAT 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    h 15.7%; Score 14.6; DB 4; Length 25; Similarity 81.0%; Pred. No. 3.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/09167681A
                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                               3.9e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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US-08-564-109C-15/c
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                                                                                                                                                                                                                                                                                                    Sequence 15, Appl. Patent No. 601810:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best
                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 904136
INFORMATION FOR SEQ ID NO:
                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                APPLICANT: Filipowicz, Witold
APPLICANT: Connelly, Sheila
TITLE OF INVENTION: Chimeric Plant Genes Possessing
TITLE OF INVENTION: Independent Regulatory Sequences
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 49
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:

APPLICATION NUMBER: WO PCT/JP95/01005
FILING DATE: 25-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                         COUNTRY: U
                                                                                                             STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: JP 6-135015
FILING DATE: 25-MAY-1994
                                                                                                STATE:
                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 31 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/01
FILING DATE: 25-JAN-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 15.7%; Score 14.6; DB 2; Length 31; Local Similarity 81.0%; Pred. No. 4.4e+03; Local Similarity 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: BENT, Stephen A REGISTRATION NUMBER: :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: JP 6-285823 FILING DATE: 26-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
SOFTWARE: PatentI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                         Application US/08564109C
                                                                            USA
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                                                                                                                                                  No. 6018103artis Corporation
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SYSTEM: PC-DOS/MS-DOS
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5504194-4/c; Patent No. 5504194
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                                                                                                                                                                       US-08-683-877-7/c
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                                                                                                                            Sequence 7, Application US/08683877 Patent No. 5776689
                                                                                                                                                                                                                                                                                                                        Matches
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              GENERAL INFORMATION:
APPLICANT: Karin, Michael
APPLICANT: Elledge, Stephen J.
APPLICANT: Aronheim, Ami
TITLE OF INVENTION: Protein Re
NUMBER OF SEQUENCES: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DI
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: S-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 800 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE: NO
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: LYMPHOCYTE ADHESION RECEPTOR FOR HIGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ST.JOHN, THOMAS P.; GALLATIN, W. MICHAEL; IDZERDA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 628,640 FILING DATE: 12-DEC-1990 FILING DATE: 12-DEC-1990 NUMBER: 325,224
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Hehman, Mary K. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
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                                                                                                                                                                                                                                                                                                                                        Match 15.7%;
Local Similarity 69.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 17-MAR-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 15-MAY-1992
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                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 33
                                                                                                                                                                                                                                                31 CGGCCGCGGCGCCAATGCATTGGGCCCCAA 3
                                                                                                                                                                                                                                                                   20 CTGCCTGGGTGACAATGCCTTGGTTTCAA 48
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                                      Protein Recruitment System
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                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                        Score 14.6; DB 6; Length 33; Pred. No. 4.5e+03;
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Pred. No. 4.5e+03;
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US-08-683-877-7
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TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 35 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION UNMER: US/08/683,877
FILING DATE: 19-UU-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Karin, Michael
APPLICANT: Elledge, Stephen J.
APPLICANT: Aronheim, Ami
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                   TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 535-9001
                                                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
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                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                            REGISTRATION NUMBER: 31 REFERENCE/DOCKET NUMBER:
                                                                                                 NAME: Campbell, Cathryn A
                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                        ZIP: 92122
                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                 TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                             San Diego
: California
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                                                                                                                                                                                                                                                                                                                                                                                             E: Campbell & Flores LLP
4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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(619) 535-8949
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                                                                                                                                                                                                                                                                               Floppy disk
                                                                                                                                                             19-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15.7%; Score 14.6; DB 1; Length 35; 69.0%; Pred. No. 4.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protein Recruitment System
                                                                                                                                                                              US/08/683,877
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                                                              P-UD 1971

 Mismatches

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US-08-431-080-9
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; Patent No. 5698686
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: KONDO, KEIJI
APPLICANT: MIURA, YUTAKA
TITLE OF INVENTION: YEAST VECTOR AND METHOD OF PRODUCING PROTEINS USING THE
TITLE OF INVENTION: SAME
FILE REFERENCE: 099441/0118
CURRENT APPLICATION NUMBER: US/09/242,690A
CURRENT FILING DATE: 1999-02-23
PRIOR APPLICATION NUMBER: PCT/JP97/02924
PRIOR PILING DATE: 1997-08-22
PRIOR FILING DATE: 1997-08-22
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PRIOR FILING DATE: 1996-08-23
NUMBER OF SEQ ID NOS: 66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                     APPLICANT: Singer, Miriam S.
TITLE OF INVENTION: Telomerase Compositions and Methods
NUMBER OF SEQUENCES: 32
                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
                                                                                                                                                    STREET:
                                                                                 COUNTRY: UI
ZIP: 77210
                                                                                                                                                 ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                                                                                                                                                    17 CAGCGAGCACTCAAATCTGCC 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33 GCAGCACCCGCGGCTGCCCCGGTGAAGAT 5
                                                                                                                                                                                                                                                                                                                                                                                                                        4 CTGGGAGCACTCCATTCTGCC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 GGAGCACTCCATTCTGCCTGGGTGACAAT 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                   Houston
                                                                                                                    TEXAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/09242690A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                              UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                           Gottschling, Daniel E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                            Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           81.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15.7%; Score 14.6; DB 4; Length 38; 81.0%; Pred. No. 4.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.6; DB 1; Length 35; Pred. No. 4.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
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US-08-185-949B-99/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 99, Application US/08185949B Patent No. 5874279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                      TELEFAX: (212) 278-05
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
TOPOLOGY: 1 MOLECULE TYPE:
                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Mark D. Cochran

APPLICANT: Richard D. Macdonald

TITLE OF INVENTION: Recombinant Infectious Bovine

TITLE OF INVENTION: Rhinotracheitis Virus
                                                                                                                                                      REGISTRATION NUMBER: ,678
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION: NAME: White, John P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: I. STREET: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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TEX: 79-0924
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                                STRANDEDNESS:
                                                                 LENGTH:
                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/185,949B FILING DATE: 03-NOV-1994
                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                          COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: Concurrently Herewith CLASSIFICATION: 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: SN 08/326,781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/431,080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63 GACTGTCGGGCCTCCAGAAAACATTGAGG 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 GACTGTCGTCCGTCAAAAAATTTCAAGG 34
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                                                                                                                                                                                                                                                                                                                                                                                              10036
                                                  nucleic acid
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                                                                 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                    E: Floppy disk
IBM 330 466 DX2
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                     linear
                                                                                                                                           (212)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   John P. White
DNA (genomic)
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                                    double
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                                                                                                                       278-0525
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0; Gaps

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US-08-185-949B-99
                                               RESULT 62
US-08-778-494B-100/c
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Patent No. 5916752
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               Sequence 100, Application US/08778494B Patent No. 5962272
                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                        Matches
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS: LENGTH: 40 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                  REPERENCE/DOCKET NUMBER: ARCD:155/PAR TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: October 20, 199
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: (FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Gottschling, Daniel E.
APPLICANT: Singer, Miriam S.
TITLE OF INVENTION: Telomerase Compositions and Methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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                                                                                                                                                                                                           Local
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                                                                                                                                                      63 GACTGTCGGGCCTCCAGAAAACATTGAGG 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               un 15.7%;
1 Similarity 69.0%;
20; Conservative
                                                                                                                      6 GACTGTCGTCCGTCAAAAAAATTTCAAGG 34
                                                                                                                                                                                        20; Conservative
                                                                                                                                                                                                           h 15.7%;
Similarity 69.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UNITED STATES OF AMERICA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Arnold, White & Durkee
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26-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SN 08/326,781
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                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                         Score 14.6;
Pred. No. 4.
                                                                                                                                                                                                           4.9e+03
                                                                                                                                                                                                                            DB 2; Length 40;
                                                                                                                                                                                            Indels
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                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                       US-08-483-511-80
                                                                                                                                                                                                                                                                                                                                      ; Sequence 80,
                                                                                                                                                                                                                                                                                                                                                                    RESULT 63
                                                                                                                                                                                                                                                                                                               ; Patent No. 6297048
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (352) 375-810
TELEFAX: (352) 372-5800
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Pace, Doran R.
REGISTRATION NUMBER: 38,261
REFERENCE/DOCKET NUMBER: CL
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 08/582,562
FILING DATE: 03-JAN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Pace, Doran R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0. CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                             APPLICANT: Jolly, Douglas J. APPLICANT: Chang, Stephen M.W. APPLICANT: Lee, William T.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Chenchik, Alex APPLICANT: Zhu, York
                                                                                                                                         ADDRESSEE: Seed and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                                 COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 03-JA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
 COMPUTER: IBM PC compatible
                                                                                                         CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                  MEDIUM TYPE:
                                                                       COUNTRY:
                                                                                        STATE:
                                                                                                                          STREET:
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                                                                                                         Seattle
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                                                                                      Washington
                                                                                                                                                                                                                                                                                                                                      Application US/08483511
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Siebert, Paul
                                                                       U.S.
                                                                                                                                                                                                                                Townsend, Kay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                             Seed and Berry
                Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA (synthetic)
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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR FULL-LENGTH CDNA TITLE OF INVENTION: CLONING
                                                   NUMBER OF SEQUENCES: 84

CORRESPONDENCES: 84
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STREET: 2421 N.W. 41st Street, Suite A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                     37 GCTGCCTGCTGCCACTCTGCGTTGTTACCACTGCTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 GCTGGGAGCACTCCATTCTGCCTGGGTGACAATGCCT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           h 15.7%; Score 14.6; DB 2; Length 40; Similarity 62.2%; Pred. No. 4.9e+03;
6300 Columbia Center, 701 Fifth Avenue
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US-09-345-294-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 9, Application US/09345294 Patent No. 6387619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                               INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 206-622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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Local Similarity 81.0%; Pred, No. 4.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 930
SLECOMMUNICATION TOTAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/483,511
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                                                                                                                                                                       NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: ARCD:155/PAR
REFERENCE/DOCKET NUMBER: ARCD:155/PAR
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/431,080 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Telomerase Compositions and Methods NUMBER OF SEQUENCES: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Gottschling, Daniel E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       72 GCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 GCCGCCGGAGAACATTGAGAT 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
SOFTMARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/345,294
FILING DATE: 30-Jun-1999
CLASSIFICATION: <Unknown>
                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Arnold, White & Durkee STREET: P.O. Box 4433
      TOPOLOGY: linear
                                                                                                                                        TELEPHONE: (512) 418-3000
TELEFAX: (713) 789-2679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: TEXAS
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                                                              LENGTH: 40 base pairs
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                                                                                                                       79-0924
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; GENERAL INFORMATION:
; APPLICANT: Buechle
                                                                                              RESULT 66
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                                                                                                                                                                                                                                                                        Query Match
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US-09-410-903-55/c ; Sequence 55, Application US/09410903
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                                     Patent No. 6420113
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                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Liebeschuetz, Joseph O. REGISTRATION NUMBER: 37,505 REFERENCE/DOCKET NUMBER: 01491
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CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
SOFTWARE: FastSE
                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 415-576-0300
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                                                                                                                                                    39 CCTGGGTTATCACAATATCGGCCATGGCTGGTTGGGC 3
                                                                                                                                                                                         37 CCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGC 73
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                                                                                                                                                                                                                                   23; Conservative
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Buechler, Joe
                                                                                                                                                                                                                                                                                                                                                                                                   43 base pairs
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Two Embarcadero Center, 8th Floor
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                                                                                                                                                                                                                                                                                                                                               linear
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                                                                                                                                                                                                                                                 15.7%; Score 14.6; DB 3; Length 43; 62.2%; Pred. No. 5.1e+03;
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                                                                                                                                                                                                                                 0; Mismatches
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Pred. No. 4.9
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US-09-410-903-55
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PRIOR TILING DATE: 1997-04-04
PRIOR PRIOR PRIOR OF THE STATE OF THE S
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Patent No. 5648211
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TITLE OF INVENTION: Chimeric Polyclonal Antibodies
FILE REFERENCE: 014907-002700US
CURRENT APPLICATION NUMBER: US/09/410,903
CURRENT FILING DATE: 1999-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: APPLICANT: APPLICANT: V
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 45 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wright, David J.
TITLE OF INVENTION: STRAND DISPLACEMENT AMPLIFICATION USING
THERMOPHILIC ENZYMES
    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                               TOPOLOGY: lir
                                                                                                                   TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Franklin Lakes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Richard J. Rodrick, Becton Dickinson and ADDRESSEE: Company
                                                                                                                                                                                                                                                                                                                         NAME: Fugit, Donna R. REGISTRATION NUMBER: 32,135
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Loca L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           37 CCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGC 73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Valkirs, Gunars
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 Becton Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Spargo, Catherine A. Walker, George T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fraiser, Melinda S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Van Cleve, Mar
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                                               linear
DNA (genomic)
                                                                                 single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US/08/229,279
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                                ; Sequence 782, Application US/09605785; Patent No. 6321716; GENERAL INFORMATION:
                                                                                           RESULT 69
US-09-605-785-782
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                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: DNA (genomic) US-08-701-269-4
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                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davin C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: INFORMATION FOR SEQ ID NO: 4:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: STRAND DISPLACEMENT TITLE OF INVENTION: THERMOPHILIC ENZYMES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: ב בכני
רדקץ: Franklin Lakes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Fugit, Donna REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                   TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Richard ADDRESSEE: Company
                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/701,269
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                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 45 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23;
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                                                                                                                                                                39 ATGCCTCCGTGTAAGCCCGAGTAGATTGGGACTATTG 3
                                                                                                                                                                                                   34 ATGCCTTGGTTTCAACACTATCGGAATGTGACTGTCG 70
                                                                                                                                                                                                                                        23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07417
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                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SD
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                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wright, David J.
VENTION: STRAND DISPLACEMENT AMPLIFICATION USING
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                                                                                                                                                                                                                                                                                                                                                 linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Donna R. 32,135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                                                                                                                                                                       15.7%; Score 14.6; DB 1; 62.2%; Pred. No. 5.2e+03;
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                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                           DB 1; Length 45;
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RESULT 70
US-08-184-009-187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/605,785
CURRENT FILING DATE: 2000-06-27
TELEX: 425066CURIMS
INFORMATION FOR SEQ ID NO: 187:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 45
TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454
                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Paoletti, Enzo
APPLICANT: Tartaglia, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY NUMBER OF SEQUENCES: 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cox, William I.
                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Cut Care Propriet 530 F1fth Avenue
                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                           CITY: New York
STATE: NY
                                                        TELEFAX:
                                                                                                                                                                                                      FILING DATE:
                                                                      TELEPHONE:
                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AGAATGCCTACCGTGCTGCAGTGCGTGAACGTGTCGGTGGTGTCT 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Retter, Marc W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Harlocker, Susan L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vedvick,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                           USA
                                                        : (212) 840-3333
(212) 840-0712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                Curtis, Morris & Safford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craig H.
                                                                                                                                                                                                    UMBER: US/08/184,009
19-JAN-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gary R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yasir A.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Thomas S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15.7%; Score 14.6; DB 4; Length 45; 57.8%; Pred. No. 5.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Robert A.
                                                                                                                454310-2530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                         ; MOLECULE TYPE: US-08-458-356-187
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US-08-458-356-187
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                                                                      Matches
                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 187, Application US/08458356
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                                                                                                                                                                                                                                     TELEX: 425066CURTMS
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 45
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 0 FILING DATE: 19-JAN-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY NUMBER OF SEQUENCES: 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Tartaglia, James APPLICANT: Cox, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: CDNA
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                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                         TELEFAX:
                                                                                                                                                                                                                                                                                                          TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                      NAME: Frommer, William S. REGISTRATION NUMBER: 25,5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE:
                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 47 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                        LENGTH:
                    35 TGCCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  35 TGCCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26;
3 TEGTTGTGTTAGTTCTCTCTAAAAATGTGGCTGCAGAGCCTGCTG 47
                                                                      26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 TGGTTGTGTTAGTTCTCTCTAAAAATGTGGCTGCAGAGCCTGCTG 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ΥY
                                                                                                                                                                                                                          47 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                    Conservative
                                                                                                                                                                                                                                                                                           : (212) 840-3333
(212) 840-0712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Paoletti, Enzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Linea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Curtis, Morris & Safford
                                                                                                                                                       CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15.7%; Score 14.6; DB 2; Length 47; 57.8%; Pred. No. 5.3e+03;
                                                                                   15.7%; Score 14.6; DB 2; Length 47; 57.8%; Pred. No. 5.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                US 08/184,009
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                                                                                                                                                                                                                                                                                                                                                            25,506
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                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                               454310-2530
                                                                    Indels
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RESULT 72 US-08-460-736-187

COUNTRY:

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RESULT 73
US-08-093-453B-7
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US-08-460-736-187
                                                                                                                                                                                                                                                                                            sequence 7, Application US/08093453B
Patent No. 5439814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 187, Application US/08460736 Patent No. 6265189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 187:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 25.506
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-333
TELEPAX: (212) 840-712
TELEY 4.2506CURTMS
FORMATION FOR ENGINEERING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 19-JAN-1994
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Curtis, Morris & Safford
STREET: 530 Fifth Avenue
                    COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS LENGTH: 47 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Tartaglia, James
APPLICANT: Cox, William I.
TITLE OF INVENTION: RECOMBINANT VIRUS IMMUNOTHERAPY
NUMBER OF SEQUENCES: 217
                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                    APPLICANT: Wang, Chin_Yen TITLE OF INVENTION: Modif NUMBER OF SEQUENCES: 13
                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0:
FILING DATE: 02-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  COMPUTER: MacIntosh
                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 57.8 es 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                      35 TGCCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCAG 79
                                                                                                                                                                                                                                                                                                                                                                                                            3 TGGTTGTGTTAGTTCTCTCTAAAAATGTGGCTGCAGAGCCTGCTG 47
                                                          30303
                                                                                                            Atlanta
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                                                                                            Georgia
                                                                                                                            191 Peachtree Street, 37th Floor
                                                                        United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
                                                                                                                                                                                                                                          Dominguez, Geraldina
                                                                                                                                                                                                                                                            Frey, Teryl K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Paoletti, Enzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                Jamie L. Greene, Jones & Askew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15.7%; Score 14.6; DB 4; Length 47; 57.8%; Pred. No. 5.3e+03;
                                                                                                                                                                                                        Modified Infectious Rubella Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US 08/184,009
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
TELEPHONE: (404) 818-: TELEFAX: (404) 818-379 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S. 07/722,334
FILING DATE: 28 JUN 1991
ATTORNEY/AGENT INFORMATION:
                                                      TELECOMMUNICATION INFORMATION:
                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Greene, Jamie L.
REGISTRATION NUMBER: 32,4
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                      SOFTWARE: Patentin Relaction CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Frey, Teryl K.
APPLICANT: Dominguez, Geraldina
APPLICANT: Wang, Chin-Yen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Greene, Jamie L.
REGISTARTION NUMBER: 32,467
REFERENCE, DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: DNA Encoding Infectious Rubella Virus NUMBER OF SEQUENCES: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE:
                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 02-JUN
                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                                                                                                        CITY: Atlanta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: 7.0
SOFTWARE: Microsoft Word
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                                                                                                                                                                                                                                                                                                                                                                                          STREET:
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                                                                                                                                                CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48 ACACTATCGGAATGTGACTGTCGGGCCTC 76
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                                                                                                                                                                                                                                                                                                                      30303
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                                                                                                                                                                                                                                                                                                                                                                                          3: Jones & Askew
191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                            PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rubella virus
                  (404) 818-3700
404) 818-3799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                404_818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO
                                                                                                                                                                  02-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA to genomic RNA
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                                                                                          32,467
                                                                            07362-0102
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Pred. No. 5.4e+03;
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9

0 Gaps

0

Length 48; Indels

Matches

COUNTRY: STATE: STREET: Query Match

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US-08-459-041A-10
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      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10, Application US/08459041A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                               TELEFAX: (404) 818-3799 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                              SUFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION NUMBER: US/08/459,041A FILING DATE: 02-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Frey, Teryl K.
APPLICANT: Dominguez, Geraldina
APPLICANT: Wang, Chin-Yen
                                                                                FEATURE:
                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 48 base pairs
                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: DNA Encoding Infectious Rubella Virus NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: NO FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                         FRAGMENT TYPE: N-terminal
                                                                                                                                                                                    HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                  NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                             OTHER INFORMATION: /note= "HindIII site"
                                                                                                                                                                                                                                                     TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
                                                                                                               LOCATION:
                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 32,467
REFERENCE/DOCKET NUMBER: 07
                                                                                                                                                                                                                                                                                                                                                                        NAME: Greene, Jamie L
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                        NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 48 base pairs
                                                                                                                                                                            -SENSE:
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                                                                                                                                                                                                                                                   nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                      Greene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       191 Peachtree Street, 37th Floor
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                                                promoter 7..23
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N: 435
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                                                                                                                                                                                                   DNA (genomic)
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                                                                                                                                                                                                                                                                                               10:
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   Score 14.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.6; DB 1; Length 48; Pred. No. 5.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
   DB 1;
Length 48;
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                                                                                                                                         Sequence 135, Application US/09038637
Patent No. 6235470
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                                                                                                      GENERAL INFORMATION:
APPLICANT: Sidran
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TELEFAX: 612-305-1228
INFORMATION FOR SEQ ID NO: 36:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
                                                                                TITLE OF INVENTION:
                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: MUETING, ANN M.
REGISTRATION NUMBER: 33,977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 228.00010201 TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 36:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: SELF-ASSEMBLING GENES, VECTORS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: NATURE TECHNOLOGY CORPORATION, ET AL.
                                                                                                                                                                                                                                                                                                         37 CCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGC 73
                                                                                                                                                                                                                                                                                                                                                  23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           48 ACACTATCGGAATGTGACTGTCGGGCCTC 76
                                                                                                                                                                                                                                                               7 CCTTGATTCCAGCAATGTTAGGACAAGGCTGGTGGGC 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/380,190A FILING DATE: 26-Aug-1999 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: MINNEAPOLIS
STATE: MINNESOTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/US98/03918 FILING DATE: 28-FEB-98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: MUETING, RAASCH & GEBHARDT, P.A. STREET: 119 NORTH FOURTH STREET, SUITE 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/09380190A
                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                    Sidransky, David
                                                                                                                                                                                                                                                                                                                                                                  15.7%; 62.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 69.0%; Pred. No. 5.4e+03;
                                                                                   DETECTION OF NEOPLASIM BY ANALYSIS OF SALIVA
                                                                 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 THEREOF
                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                  Score 14.6; DB 4; Length 48; Pred. No. 5.4e+03;
                                                                                                                                                                                                                                                                                                                                                  14; Indels
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                                                                                                                                                                                                                                                                                                                                            Gaps
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; MOLECULE TYPE: Genomic DNA US-09-038-637-135
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/07613083B Patent No. 5340727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 619/678-5099
INFORMATION FOR SEQ ID NO: 135:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for CURRENT APPLICATION DATA:
CLASSIFICATION: 435
PRIOR APPLICATION DATA: This appl. is a c-i-p of APPLICATION NUMBER: U.S. 07/470,674
                                                     SOFTWARE: WordPerfect 5.1 conv. to ASCII CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/613,083B
FILING DATE: 19911114
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/15
FILING DATE: 12-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: GPID' Fragments and RecTITLE OF INVENTION: DNA Expression Vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: U0/1/1/2 APPLICATION NUMBER: 28-DEC-1995 O8/152,313
                                                                                                                  COMPUTER: AST BIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                      CITY: La Jolla
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/038,637 FILING DATE: 10-MAR-1998
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                                                                                                                                                                 MEDIUM TYPE:
                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                             ADDRESSEE: Scripps Clinic and Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                                                                                                                      92037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 15.5%; Score 14.4; DB 4; Similarity 93.8%; Pred. No. 4.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Haile, Lisa A.
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                                                                                                                                                                                                                                                                         10666 No. 5340727th Torrey Pines Road
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619/678-5099
                                                                                                                                                                                                                      USA
                                                                                                                                                                                                                                                                                                                                                                                                                 Ruggeri, Zaverio M.
Ware, Jerry, inventors
on behalf of Scripps Clinic and Research
                                                                                                                          E: Diskette-5.25 inch, 1.2 Mb
AST Bravo IBM PC comp. (386SX)
SYSTEM: MS DOS version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                      Foundation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 4; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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; TOPOLOGY: Linear US-07-613-083B-2
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US-08-474-021-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/08474021 Patent No. 5650282
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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TELEFAX: (215) 923-2189
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                 HYPOTHETICAL: NO ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                          TELEFAX: 202-962-8300
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Keating, Mark T.
APPLICANT: Leppert, Mark F.
APPLICANT: Morris, Colleen A.
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF SUPRAVALVULAR
TITLE OF INVENTION: AORTIC STENOSIS AND WILLIAMS SYNDROME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 04-Jan-1990 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 19: TELECOMMUNICATION INFORMATION: TELEPHONE: 202-962-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                          MOLECULE TYPE:
                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Barron, Alexis
REGISTRATION NUMBER: 22,
REFERENCE/DOCKET NUMBER:
                                                                                                           TOPOLOGY:
                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                     NAME: Ihnen, Jeffre REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                  ORGANISM:
                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1201 New York Ave., N.W., Suite 1000
                                                                                                                                                                       32 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                            202-962-8300
                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Venable, Baetjer, Howard & Civiletti
                                                                            NO
                                                                                                                                                                                                                                                                                                                   Jeffrey L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Floppy disk
                                                                                            DNA (genomic)
                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single stranded
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                                                                                                                                                                                                                                                                                                                                                                                 US/08/041,576
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                                                                                                                                                                                                                                                                                                       28,957
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US-08-474-020A-2/c
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                                                                                                          밁
                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens US-08-474-020A-2
                                 PCT-US94-03426-2/c
                                                     RESULT 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/08474020A Patent No. 5840489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
Sequence 2, Application PC/TUS9403426 GENERAL INFORMATION:
                                                                                                                                                                           Matches
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                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 202-962-8300 INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Keating, Mark T.
APPLICANT: Leppert, Mark F.
APPLICANT: Morris, Colleen A.
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF SUPRAVALVULAR
TITLE OF INVENTION: AORTIC STENOSIS AND WILLIAMS SYNDROME
                                                                                                                                                                                                                                                                               ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION TELEPHONE: 202-962-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 08/041,576
FILING DATE: 05-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Inhen, Jeffrey L.
REGISTRATION NUMBER: 26,957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                              ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 202-962-8300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 19
TLECOMMUNICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH:
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                                                                                                      31 AAACTGGCTCCCAGGCCTCCAGAA 8
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                                                                                                                                                                                                                                                                                                                                                                                             32 base pairs
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                                                                                                                                                                                                                                                                                                                                                linear
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                                                                                                                                                                                                                                                                                                               NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Floppy disk
                                                                                                                                                                                                                                                                                                                              DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15.5%;
75.0%;
                                                                                                                                                                                        15.5%; Score 14.4; DB 2; Length 32; 75.0%; Pred. No. 5.4e+03;
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                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19780-105509
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                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Indels
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RESULT 82
US-08-840-466A-10/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Keating, Mark T.
APPLICANT: Leppert, Mark F.
APPLICANT: Leppert, Mark F.
APPLICANT: Morris, Colleen A.
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF SUPRAVALVULAR
TITLE OF INVENTION: STENOSIS AND WILLIAMS SYNDROME
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
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NAME: Ihnen, Jeffrey L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 32 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORIGINAL SOURCE:
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ANTI-SENSE: N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
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                                                                                                                                                                                                   MCKee, Marian L.
O'Brien, Alison D.
O'Brien, Alison D.
Wachtel, Marian R.

TITLE OF INVENTION: Method Of Stimulating An Immune Response
By Administration Of Host Organisms That Express Intimin
Alone Or As A Fusion Protein With One Or More Other
ZIP: 20005-3315
COMPUTER READABLE FORM:
                                                                                                                                                              NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                APPLICANT: Stewart, C. Neal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31 AAACTGGCTCCCAGGCCTCCAGAA 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   58 AATGTGACTGTCGGGCCTCCAGAA 81
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Similarity 75.0%;
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                     COUNTRY: USA
ZIP: 20005-3315
                                                           STATE: D.O
                                                                            CITY: Washington
                                                                                                  STREET: 1300 I Street, N.W., Suite 700
                                                                                                                                      ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Venable, Baetjer, Howard & Civiletti
1201 New York Ave., N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA (genomic)
                                                                                                                        Dunner, L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCT/US94/03426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28,957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19780-105509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.4; DB 5; Length 32; Pred. No. 5.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 83
US-09-696-188B-10/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6406885
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10, Application US/09696188B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15.5%; Score 14.4; DB 4; Length 34; Local Similarity 75.0%; Pred. No. 5.5e+n3.
                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           O'Brien, Alison D.
Wachtel, Marian R.
Warthel, Marian R.
FITLE OF INVENTION: Method of Stimulating An Immune Response
By Administration Of Host Organisms That Express Intimin
Alone Or As A Fusion Protein With One Or More Other
TELECOMMUNICATION INFORMATION:
                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Stewart, C. Neal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36 GCCTTGGTTTCAACACTATCGGAA 59
                                                                                                                                                                                APPLICATION NUMBER: US/09/696,188B FILING DATE: 26-Oct-2000 CLASSIFICATION: <Unknown>
                 NAME: Boone, Laural S.
REGISTRATION NUMBER: 43,505
REFERENCE/DOCKET NUMBER: 04995.0029-00000
                                                                                                                  APPLICATION NUMBER: 08/840,466 FILING DATE: 1997-04-18
                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 1300 I Street, N.W., Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 20005-3315
                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 34 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/840,466A FILING DATE: 18-Apr-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (202) 408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 43,505
REFERENCE/DOCKET NUMBER: 04995.0029-00000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Boone, Laural S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mckee, Marian L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dunner, L.L.P.
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US-09-375-318-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-375-318-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 6, Application US/09375318; Patent No. 6468791
GENERAL INFORMATION:
                                         Matches
                                                                               Query Match
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                                                                                                                                                                                                                                                            TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
                                                           Local
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                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 33,287
REFERENCE/DOCKET NUMBER: 92
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/375,318
FILING DATE: 16-Aug-1999
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Galas, David {\mathfrak J}. TITLE OF INVENTION: CHROMOSOME 1 GENE AND GENE PRODUCTS RELATED TO
                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Tanzi, Rudolph E.
22 GCCTGGGTGACAATGC 37
                                       15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    36 GCCTTGGTTTCAACACTATCGGAA 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25 GCCTTGGTTTGATCAGGATCCGTA 2
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                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 15.5%;
Similarity 75.0%;
                                                                                                                                                                           LENGTH: 39 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Seattle
STATE: Washington
                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: SEED and BEERY LLP STREET: 701 Fifth Ave, Suite 6300
                                                                                                                                                                                                                                                                                                                                                                                       NAME: Verna, James M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 98104-7092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 34 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
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Bird, Thomas D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wasco, Wilma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schellenberg, Gerard D.
                                                                                                                                                                linear
                                                    15.5%; Score 14.4; DB 4; Length 39; 93.8%; Pred. No. 5.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ALZHEIMER'S DISEASE
                                     0;

 Mismatches

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                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                920010.571C1
                                     Indels
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                                     0
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В

23 GCCTGGGTGACAAAGC 38

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Sequence 19, Application US/08687559
PATENT NO. 5955647
GENERAL INFORMATION:
APPLICANT: Fitchen, John H.
APPLICANT: Beachy, Roger N.
TITLE OF INVENTION: METHOD FOR USING TOBACCO MOSAIC VIRUS TO TITLE OF INVENTION: OVERPRODUCE PEPTIDES AND PROTEINS
                                                                                                                                                                                                    밁
                                                                                                                                  US-08-687-559-19/c
                                                                                                                                                                                                                                                                                                                                     US-08-301-722A-6
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US-08-301-722A-6/c
                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 6, Application US/08301722A Patent No. 5756307
                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTMARE: Patentin Release #1.0, VOURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/301,722A
FILING DATE: 07-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Uhl, George R.
APPLICANT: Vandenbergh, David
APPLICANT: Persico, Antonio
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 11:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Murphy Jr., Gerald M. REGISTRATION NUMBER: 28,977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
                                                                                                                                                                                                                                                                                   ocal
                                                                                                                                                                                                 39 TGGGGGCCCTGCATGCGTCCTGGG 16
                                                                                                                                                                                                                                                                   18;
                                                                                                                                                                                                                               5 TGGGAGCACTCCATTCTGCCTGGG 28
                                                                                                                                                                                                                                                                               15.5%; Score 14.4;
Similarity 75.0%; Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22040-3487
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Falls Church
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 40 base pairs
                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    YES
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                                                                                                                                                                                                                                                                                                                                   /label= consensus
/note= "consensus sequence of VNTR element in 3'
untranslated region of HUDAT cDNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA
                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                               6e+03;
                                                                                                                                                                                                                                                                                          DB 1; Length 40;
                                                                                                                                                                                                                                                             6; Indels
                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                             0,
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                                                                                                                                                                                             STATE:
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Sequence 44, Application US/08145704 Patent No. 5567604
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 619-678-5099
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                     APPLICANT: Fennewald, Susan
APPLICANT: Zendegui, Joseph G.
APPLICANT: Joshua O. Ojwang
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
NUMBER OF SEQUENCES: 45
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rando, Robert F.
APPLICANT: Fennewald, Susan
APPLICANT: Zendegui, Joseph
APPLICANT: Joshua O. Ojwang
                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: Genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 619-678-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                      COUNTRY: U.S.A. ZIP: 77010-3095
                                  APPLICATION NUMBER: US/08/145,704 FILING DATE: 28-OCT-1993
                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Learn, June M. REGISTRATION NUMBER:
                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: PCT/US95/01467 FILING DATE: 03-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: No. 5955647ember 18, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         37 TTGGACCTCTAGAATACATATAGG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            68 TCGGGCCTCCAGAAAACATTGAGG 91
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                                                                                                                                                                                                                                               Houston
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                                                                                                                                                                                                                                                          E: Fulbright & Jaworski
1301 McKinney, Suite 5100
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75.0%;
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Pred. No. 6.2e+03;
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                                   INFORMATION FOR SEQ ID NO:
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LENGTH: 47 base pairs
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REFERENCE/DOCKET NUMBER: D.STELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5151
                                                                                                                                         APPLICATION NUMBER: US 08/053,027
FILING DATE: 23-APR.1993
ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REGISTRATION NUMBER: 32,714
                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
NUMBER OF SEQUENCES: 52
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NAME: Paul, Thomas D.
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                   SEQUENCE CHARACTERISTICS
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                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 713/651-5151
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TOPOLOGY: 11r
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FILING DATE: 28-OCT-1993
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                                                         TELEX:
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                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
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LENGTH: 47 base pairs
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                                                                                                                               REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
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Local Similarity 65.6%;
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1301 McKinney, Suite 5100
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Hogan, Michael E.
                                                                         713/651-5246
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Zendegui, Joseph G.
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                                                                                                                                                                                                                                                                                                                                                                       Release #1.0, Version #1.25
                                                                                                                                                                                                                                                           PCT/US94/04529
                                                                                                                                                                                                                                                                                                                                     US/08/987,574
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                                                                                                                                 D-5574-CIP
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Pred. No. 6.4e+03;
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                                                                                                                                      ; MOLECULE TYPE: US-08-535-168-44
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                                                                   Matches
                                                                                      Query Match
Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08 FILING DATE: 23-APR-1993 ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
NUMBER OF SEQUENCES: 52
                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
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                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US94/04529 FILING DATE: 28-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP:
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                                                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                                                                                              TELEPHONE: 713/651-5151
TELEFAX: 713/651-5246
                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 32.
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A. ZIP: 77010-3095
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                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                    Local
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                                 43 TTTCAACACTATCGGAATGTGACTGTCGGGCC 74
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6 TTGGGAAACCCTTGGAACCTGACTGGCC 37
                                                                                  h 15.5%; Score 14.4; DB 4; Similarity 65.6%; Pred. No. 6.4e+03;
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Zendegui, Joseph G.
Ojwang, Joshua O.
Hogan, Michael E.
                                                                                                                                                                                                                       47 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Fulbright & Jaworski
1301 McKinney, Suite 5100
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rando, Robert F
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                                                                                                                                                                                                                                                                                                                                                                                   Thomas D.
                                                                                                                                                       DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                32,714
                                                                   0; Mismatches
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                                                                                                     Length 47;
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; Sequence 44, Application US/08682255A
; Patent No. 6323185
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US-09-017-974-44
                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 44:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Rando, Robert F.
APPLICANT: Ojwang, Joshua O.
APPLICANT: Hogan, Michael E.
APPLICANT: Wallace, Thomas L.
APPLICANT: Cossum, Paul A.
TITLE OF INVENTION: Artiviral (
TITLE OF INVENTION: Tetrad Form.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                APPLICANT:
APPLICANT:
APPLICANT:
                                                                                     APPLICANT: APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 47 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS Word 97 (saved as .txt file)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 09-DEC-97
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 713/238-8008
                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 33,962
REFERENCE/DOCKET NUMBER: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/037,374 FILING DATE: 04-FEB-97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/017,974
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                                                                                                                                                                                                                                                                            43 TTTCAACACTATCGGAATGTGACTGTCGGGCC 74
                                                                                                                                                                                                                                                                                                                    21;
                                                                                                                                                                                                                                          6 TTGGGAAACCCTTGGAACCTGACTGGCC 37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                McDaniel, C. Steven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Texas
                                                                                                                                                                                                                                                                                                                    Conservative
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                                Ojwang, Joshua O.
Hogan, Michael E.
Mazumder, Abhijit
                                                                                                      Rando, Robert F.
                                                                Zendegui, Joseph G.
                                                                                   Fennewald, Susan
               Pommier, Eyves
                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                           15.5%; Score 14.4; DB 4; Length 47; 65.6%; Pred. No. 6.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-Viral Guanosine-Rich
Tetrad Forming Oligonucleotides
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; Patent No. 6355785 ; GENERAL INFORMATION:
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US-09-429-130-44
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                                                                                                                                                                                                       Sequence 44, Application US/09429130 Patent No. 6355785
                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 713/238-8008 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Anti-vitation TITLE OF INVENTION: Oligonucleotides -- ~~ cromences: 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: McDaniel, C. Steven REGISTRATION NUMBER: 33,9 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: /13/238-8008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 25-MARCH-96
APPLICATION NUMBER: 60/C
FILING DATE: 19-MARCH-96
                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/0:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US FILING DATE: 23-OCT-95 APPLICATION NUMBER: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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              TITLE OF INVENTION: Anti-Viral Guanosine-Rich
Oligonucleotides
NUMBER OF SEQUENCES: 87
                                                                                                                                                                    APPLICANT: Rando, Robert F.
                                                                                                                                                                                                                                                                                                                                        43 TTTCAACACTATCGGAATGTGACTGTCGGGCC 74
                                                                                                                                                                                                                                                                                                                                                                          21;
                                                                                                                                                                                                                                                                                                       6 TTGGGAAACCCTTGGAACCTGACTGGCC 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid
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                                                                                                  Ojwang, Joshua O.
Hogan, Michael E.
                                                                                  Pommier, Eyves
                                                                Mazumder, Abhijit
                                                                                                                                              Fennewald, Susan
                                                                                                                                  Zendegui, Joseph G.
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N: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                     15.5%; Score 14.4; DB 4; Length 47; 65.6%; Pred. No. 6.4e+03; ative 0; Mismatches 11; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-Viral Guanosine-Rich
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US-09-429-130-44
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             APPLICANT: Zendegui, Joseph G.
APPLICANT: Ojwang, Joshua O.
APPLICANT: Hogan, Michael E.
APPLICANT: Hogan, Michael E.
APPLICANT: Mazumder, Eyves
APPLICANT: Mazumder, Abhiljit
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
NUMBER OF SEQUENCES: 52
                                                                                                                                                                                         APPLICANT: Rando, Robert F. APPLICANT: Fennewald, Susan APPLICANT: Zendegui, Joseph
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 15.5%; Score 14.4; DB 4; Length 47; Local Similarity 65.6%; Pred. No. 6.4e+03; les 21; Conservative 0; Mismatches 11; Indels
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SOFTWARE: MS WORD 97 (saved as .txt file)
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/238-8010
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   43 TTTCAACACTATCGGAATGTGACTGTCGGGCC 74
                                                                                                                                                                                                                                                                                                                                                                                                                     6 TTGGGAAACCCTTGGAACCTGACTGGCC 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/014,007 FILING DATE: 25-MARCH-96 APPLICATION NUMBER: 60/013,688 FILING DATE: 19-MARCH-96 APPLICATION NUMBER: 60/016,271 FILING DATE: 17-APRIL-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: McDaniel, C. Steven REGISTRATION NUMBER: 33,962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/001,505 FILING DATE: 19-JULY-95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/682,255 FILING DATE: <Unknown>
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STATE: Texas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: <Unknown>
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                                                                                                                                                                                                            Fennewald, Susan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 47 base pairs
                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOSTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
PRIOR APPLICATION DATA:
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NAME: McDaniel, C. Steve
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: NOVEL HUMAN GROWTH FACTORS
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: YU, GUO-LIANG APPLICANT: DILLON, PATRI
                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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FILING DATE: 23-027-95; 17-APRIL-96; 25-MARCH-96; 19-MARCH-96; 23-FILING DATE: APRIL-96; 17-APRIL-96
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                   FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                     STREET: LIUU NEW CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: McDaniel, C. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A. ZIP: 77002-2912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                            APPLICATION NUMBER: US/08/916,576B
                                                                                                                                                                                                                                            STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US96/11786
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                                                                                                                                                       MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 15.5%; Local Similarity 65.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  43 TTTCAACACTATCGGAATGTGACTGTCGGGCC 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 TTGGGAAACCCTTGGAACCTGACTGGCC 37
                                                                                                                                                                                                   20005-3934
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                                                                                                                                                                                                                                            엉
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                                                                                                                                                                                                                                                                                 E: STERNE, KESSLER, GOLDSTEIN & 1100 NEW YORK AVENUE, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Conley, Rose & Tayon, P.C. 600 Travis, Suite 1850
                                                                                                                                                                                                                                                                                                                                                                                                    DILLON, PATRICK J.
EBNER, REINHARD
                                                                                                                                                                                                                      SD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA (genomic)
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3ER: 33,962
                                                                                                                                                                                                                                                                                                      KESSLER, GOLDSTEIN & FOX, P.L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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Pred. No. 6.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 47;
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US-09-049-475-7
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                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FASTSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/049,475
FILING DATE: 27-MAR-1998
CLASSIFICATION: 800
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ADDRESSEE: PENNIE & EDMONDS LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: CDN
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 93
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)7909090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9
FILING DATE: 27-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: O'Reilly, David APPLICANT: Thomas, Christopher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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ORIGINAL SOURCE:
                                     ANTI-SENSE:
                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: single
                                                                                                        STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                           .H. 7.1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                               TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       53 ATCGGAATGTGACTGTCGGGCCTCCAGAAAACATTGAGGT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41 ATAAGAATGTGCCCCTTCTTTCATCAGAGAAGCTTGCGGT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24; Conservative
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                                                                                                                                                                                   nucleic acid
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COTUZZÍ, LAUTA A
ORIGINAL A
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RESULT 96
US-08-513-974B-81/c
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APPLICANT: FUJii, Ryo
APPLICANT: Ohtaki, Tetsuya
APPLICANT: Fukusumi, Shoji
APPLICANT: Ohqi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL
                                                       APPLICATION NUMBER: JP 6
FILING DATE: 11-AUG-1994
PRIOR APPLICATION DATA:
                                                                                                             APPLICATION NUMBER: JP 6
FILING DATE: 30-SEP-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-270017
FILING DATE: 02-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/
FILING DATE: 10-AUG-1995
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: DIKE, BRONSTEIN, ROBERTS & CUSHMAN, LLP
APPLICATION NUMBER: JP 6 FILING DATE: 11-AUG-1945 PRIOR APPLICATION DATA:
                                                                                                                                                                       FILING DATE: 30-SEP-PRIOR APPLICATION DATA:
                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 20-JAN
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 16-MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 19-AUG
                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/513,974B FILING DATE: 14-SEP-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GAACTGGGACCACTCCAGT 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                         30-SEP-1994
                                                                                                                                                                                                                                                                                                           28-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                  20-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                          16-MAR-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      380
                                                                                                                                                        JP 6-236356
                                                                                                                                                                                                                                                                                                                                                                                     JP 7-007177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Release #1.0, Version #1.30
                                                                                                JP 6-189274
                                                                                                                                                                                                                JP 6-236357
                                                                                                                                                                                                                                                                                                                             JP 6-326611
                                                                                                                                                                                                                                                                                                                                                                                                                                           JP 7-057186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP 7-093989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCT/JP95/01599
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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; MOLECULE TYPE: cDNA US-08-340-011-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-513-974B-81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 16, Application US/08340011 Patent No. 5776755
 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                   Query Match
                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/959

FILING DATE: 09-OCT-1992

ATTORNEY/AGENT INFORMATION:

NAME: MEYERS, Thomas C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 81: SEQUENCE CHARACTERISTICS: LENGTH: 27 base pairs
                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 32
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Alitalo, et al.
TITLE OF INVENTION: FIT4, A NOVEL RECEPTOR TYROSIN KINASE
NUMBER OF SEQUENCES: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: JP 6-
FILING DATE: 11-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Resnick, David S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 45
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-523-3400
                                                                                                                                                                                TELEFAX: 25-3856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                         TOPOLOGY:
                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                     TELEPHONE: 312/474-0448
                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/340,011 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Resnick, David S
REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 15.3%; Score 14.2; DB 3; Length 27; Local Similarity 84.2%; Pred. No. 6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26 CTGTAGGGCATCCATAAAA 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65 CTGTCGGGCCTCCAGAAAA 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60606-6402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  617-523-6440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   United States of America
                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Marshall, O'Toole, Gerstein, Murray & Borun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA
                                                                                                         single
 15.3%;
                                                                                                                                                                                                                                                                                                                                                                07/959,951
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                                                                                                                                                                                                                                                                                         36,989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
Score 14.2; DB 1;
Pred. No. 6.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
               Length 28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-901-710-16
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                                                                    Matches
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                                                                                                       Query Match
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                                                                                                                                                                                                                                                                  TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Alitalo, Kari
APPLICANT: Aprelikova, Olo
APPLICANT: Pajusola, Katri
APPLICANT: Armstrong, Elin
APPLICANT: Korhonen, Jaana
                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/257,754
FILING DATE: 09-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/959,951
FILING DATE: 09-OCT-1992
                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/3.
FILING DATE: 14-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: FLT4, A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Pain
                                                                                                                                                                                                                            LENGTH:
                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: sing
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CITY: Chicago
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TELEFAX: 312/474-0448
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                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 38,153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                                                                        local
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2 GGGATCCCTCCATGCTGCC 20
                                  6 GGGAGCACTCCATTCTGCC 24
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                                                                                    h 15.3%; Score 14.2; DB 3; Length 28; Similarity 84.2%; Pred. No. 6.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RY: United States of America 60606-6402
                                                                                                                                                                                                                                                                                                                                                                                     Gass, David A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/08901710
                                                                                                                                                                                                                               28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kaipainen, Arja
Matikainen, Marja-Terttu
IVENTION: FLT4, A RECEPTO
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Armstrong, Elina
Korhonen, Jaana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pajusola, Katri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aprelikova, Olga
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Marshall, O'Toole, Gerstein, Murray & Borun
                                                                                                                                                                                           single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08/340,011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US/08/901,710
                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A RECEPTOR TYROSINE KINASE, AND USES

 Mismatches

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                                                                    Indels
                                                                    0;
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                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                    0
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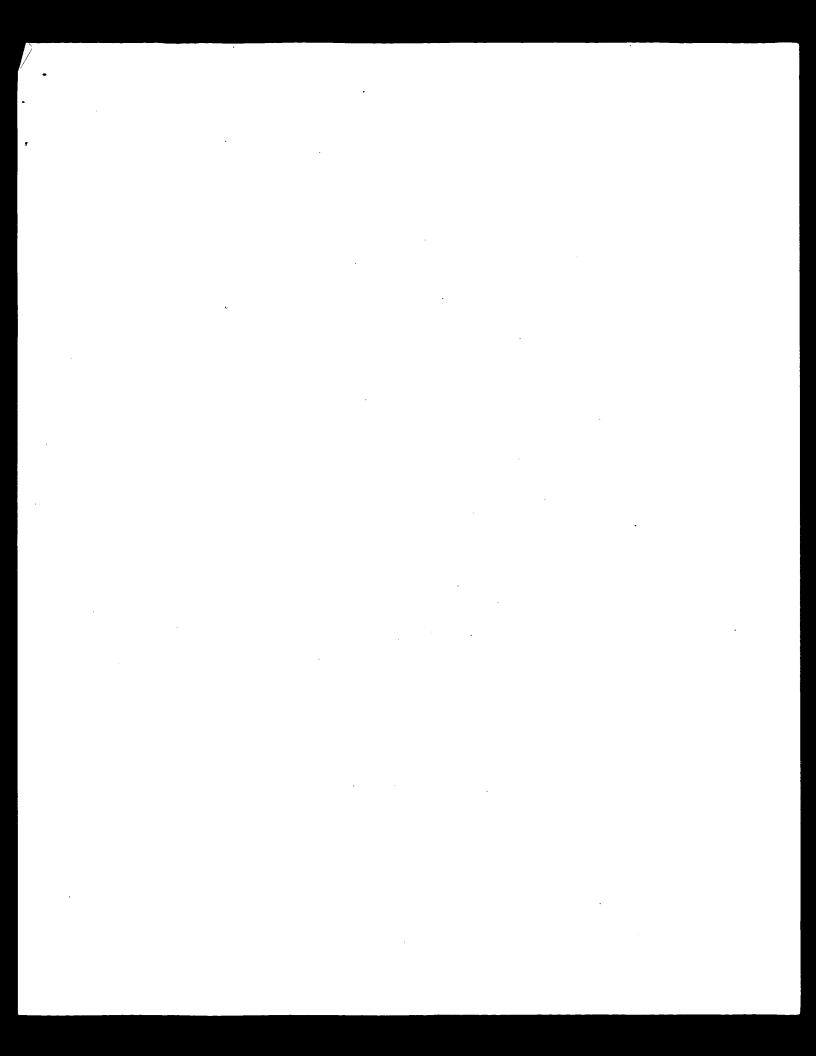
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                                                                                                                                                                                                                                                                                                                                                                       US-08-627-907A-8
                                                                                                                                                                                                                                                                                                                                                                                            RESULT 100
                                                                                                                                                                                                                                                                                                                                Sequence 8, Application US/08627907A Patent No. 6060302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 16, Application US/08331081B Patent No. 5998697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: HIRANO, Naoto
APPLICANT: HIRAI, Hisamaru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (613) 234-5233
INFORMATION FOR SEQ ID NO: :
                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: Canada 2,126,138
-FILING DATE: June 17, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC or Compatibles
OPERATING SYSTEM: PC Dos 5.0
SOFTWARE: Wordperfect 5.1 (Patseq.txt)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 234-1907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: McFadden, Fincham
STREET: 225 Metcalfe Street, Suite 606
                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                               TITLE OF INVENTION: HUMAN PHOSPHOLIPASE C-ALPHA AND DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Devlin, Robert H. TITLE OF INVENTION: Transger
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                        STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 26 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/331,081B FILING DATE: October 20, 1994
                                                                                                  COUNTRY:
                                                                                                                   STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Diskette 3.5 inch, 1.44 Mb MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 15.3%;
Local Similarity 70.4%;
les 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 ATTCTGCCTGGGTGACAATGCCTTGGT 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                 28 ATTCGGCCCGGGTGGCCAAGCTTGGAT 2
                                                                                                                                   New York
                                                                                 10020
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fincham, H. Ian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ontario
                                                                                                                   New York
                                                                                                                                                        1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29 base pairs
                                                                                                  USA
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    Mismatches

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Search completed: July 5, 2003, 13:10:03 Job time: 80 secs
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                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nu
                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: SH
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
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APPLICATION NUMBER: JP 5-238402
FILING DATE: 24-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: PCT/
FILING DATE: 22-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                   DESCRIPTION: /desc = "mutagenic primer"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/627,907A
                                                                                                                                                                                     Local
                                                                                                                    37 CCTTGGTTTCAACACTATCGGAATGTG 63
                                                                                                                                                               il Similarity
19; Conserv
                                                                                4 CCTTGGTCTGGTCACTCTAAGAATCTG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HALEY Jr., James F.
                                                                                                                                                                 Conservative
                                                                                                                                                          15.3%; Score 14.2; DB 3; Length 30; 70.4%; Pred. No. 6.3e+03; ative 0; Mismatches 8; Indels
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Result
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Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Dunn,D., Aoyaqi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                    Mus musculus
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clone UUGC1M0361N06 F, DNA sequence.
                                       Mouse whole genome scaffolding with paired end reads from 10kb
                                                and Wright, D., Weiss, R.
                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Rosso, M., Strizhov, N., Li, Y., Reiss, B., Dekker, K. and Weisshaar, B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat)
                                                                                 Unpublished
                                                                                                                                     A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA
                                                                                                                                                                                                                                                                                                                                                        Arabidopsis thaliana
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Insert Length: 10000 Std Error:
Plate: 0361 row: N column: 06
Seg primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gli4732114/gb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and musified means to be complementary to the insert adaptors and musified means but has appealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005\ \mathrm{inch} orifice at constant velocity. The sheared DNA
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                                                        4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan Email: ysuzukiéims.u-tokyo.ac. p Suzuki, Y. Yoshitomo-Wakagawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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                                                                                                                                                                                    Department of Virology Institute of Medical Sc
                                                                                                                                                                                                                                                                                   mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
                                                                                                                                                                                                                                                                                                                                 y. Y., Nakamura, Y., Suyama, A. and Sugano, S. Diverse transcriptional initiation revea
                                                                                                                                                                                                                                                                                                                                                                   Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha, Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AU106864 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8 AAAATCCGCAGTTTTCTACATTAATGGATTGTTTCTGTC 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             availability can be found at: http://www.mpiz-koeln.mpg.de/GABI-Kat/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut fuer
                                                                                                                                                                                                                                          Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
1 (bases 1 to 50)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AU106864.1 GI:13556385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AU106864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NBLAN21NR1, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          were directly sequenced to determine the genomic sequence flanking the insertion. Sequences displaying significant similarity to the A. thalaina nuclear genome sequence were processed for submission. T-DNA derived sequences were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /Clone_lib="Arabidopsis thaliana T-DNA insertion lines" /note="PCR was performed on DNA from Arabidopsis thaliana plants (T1) which were transformed with the T-DNA from
/organism="Homo sapiens"
                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               removed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         insertions. The DNA fragment(s) resulting from the PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vector pAC161. The lines contain one or more T-DNA
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/db_xref="taxon:3702"
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                                                                                                                                                                                      Science, University of Tokyo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 18.2; DB 17; Length 49; Pred. No. 2.7e+04;
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                                                                                                                                                                                                                                                                                                                               initiation revealed by fine, large-scale
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                  DEFINITION
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                                                             AZ317115/c
                                                                                       RESULT 5
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44 AATACCCTGGTGGCACTACGGGCGGAGTGTGCCTTTAATCCCT 2
                                                                                                                                                                                                                                                         33 AATGCCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCT 75
1M0035E20R Mouse 10kb plasmid \bar{U}UGC1M library Mus musculus genomic clone UUGC1M0035E20\ R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               38 GCATTCCAACCAGCCTGGGTGACA 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to give a tight size distribution ( 4\ kb ). The v+i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned populat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TA287A02Q 44 bp DNA linear GSS 13-DE T. brucei sheared genomic DNA clone 287a02, reverse sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Trypanosoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Trypanosoma brucei.
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Similarity 83.3%;
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                                                                                                                                                                                                                                                                                                                  Conservative
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/db_xref="taxon:5691"
/clone="287a02"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Trypanosoma brucei"
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                                                                                                                                                                                                                                                                                                                                              18.7%;
                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                              Score 17.4; DB 17; Length 44; Pred. No. 5e+04;
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Pred. No. 4.
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                                                            46 bp
                                                                                                                                                                                                                                                                                                                                        5e+04;
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                                                                                                                                                                                                                                                                                                            16; Indels
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                                                            GSS 29-SEP-2000
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                           Match 18.7%;
Local Similarity 77.8%;
                                                                                                                                                                                                             38 TGGGCAGTCTCCATTCTGACTGGGAGA 12
H46101 39 bp mRNA linear EST 31-JUL-
y017d02.rl Soares adult brain N2b5HB55Y Homo sapiens cDNA clone
IMAGE:178179 5' similar to gb:D14531 60S RIBOSOWAL PROTEIN L9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                      5 TGGGAGCACTCCATTCTGCCTGGGTGA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0035 row: E column: 20
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AFT29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant,
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                                                                                                                                                                                                                                                                                                                                                           Pred. No. 5.1e+04;
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  12 GCAATCAGACTGTCGACATTCCAGAAAA 39
                                                  56 GGAATGTGACTGTCGGGCCTCCAGAAAA 83
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1 (bases 1 to 39)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Warra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R.
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High quality sequence starts: 1
High quality sequence stops: 1
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Contact: Wilson RK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Source: IMAGE Consortium, LLNL free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Washington University School of Medicine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: est@watson.wustl.edu
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                                                                                                                                      Similarity
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                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                    double-stranded CDNA was size selected, ligated to Eco RI adapters (pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT713 vector (pharmacia), Library went through one round of normalization to a Cot = 53. Library constructed by Bento Soares and M.Fatima Bonaldo. The adult brain RNA was provided by Dr. Donald H. Gilden. Tissue was acquired 17-18 hours after death which occurred in consequence of a ruptured aortic aneurysm. RNA was prepared from a pool of tissues representing the following areas of the brain: frontal, parietal, temporal and occipital cortex from the left and right hemispheres, subcortical white matter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /dev_stage="55-year old"
/lab_host="pHIOB (ampicillin resistant)"
/lab_host="pHIOB (ampicillin resistant)"
/note-"Organ: brain; Vector: pT7T3D (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5'
                                                                                                                                                                                                                                                                             medulla.
                                                                                                                                                                                                                                                                                                     basal ganglia, thalamus, cerebellum, midbrain, pons and
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/clone="IMAGE:178179"
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/db_xref="GDB:3840375"
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Std Error: 0.00
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Pred. No. 7.8e+04;
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COMMENT
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                                                   University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                  Unpublished (2000)
                                                                                                                                                                                                                                Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                      Contact: Robert B. Weiss
                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ659644 39 bp DNA linear GSS 14-DEC-20
1M0537G10F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0537G10 F, DNA sequence.
                                                                                                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 39)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jag
Email: ysuzuki@ims.u-tokyo.ac.j
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites EMBO REP. 2 (5), 388-393 (2001)
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AU103467
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/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
a 9 c 11 g 15 t
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Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan Email: ysuzuki@ims.u-tokyo.ac.jp Suzuki,r., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        58 AATGTGACTGTCGGGCCTCCAGAAAACATTGA 89
                                                                                                                                                                                                                                                                             Y., Nakamura, Y., Suyama, A. and Sugano, S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
                                                                                                                                                                                                                                                                                                                                                             Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J.,
H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okub
                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AU106000 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
                                                                                                                                                              Department of Virology
                                                                                                                                                                                        Contact: Yutaka Suzuki
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HEP00195, mRNA sequence.
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Insert Length: 10000 Std Erro
Plate: 0537 row: G column: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwMp4 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coil XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                              2 (5), 388-393 (2001)
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/sex="Male"
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Pred. No. 1.5e+05;
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                                                                                                                                                                            38 CTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCC 77
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 50)
Suzuki, 7. Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha.
J., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki
Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
AU104835 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                      4 CTTGGGTCCGTCGCTTCGGTGTCCCTGTCGGGCTTCC 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ysuzuki@ims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
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/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
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/db_xref="taxon:9606"
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Pred. No. 1.7e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 CCTTCTGCCTAGGTCCCAACGCTTCGG 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
S. Construction and characterization of a full length-enriched and
a 5'-end-enriched CDNA, library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
                                             Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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AU104835
                                                                                                                                                                                                                                                         Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn, J., Zimmerman,J., and Ecker,J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SALK_084858.34.85.x Arabidopsis thaliana TDNA insertion
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                                                                                                                                                                                       Arabidopsis Genome Unpublished (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BH855636
                            Email: ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BH855636.1 GI:21705226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis thaliana genomic clone SALK_084858.34.85.x, DNA
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This is single pass sequence recovered from the left border of
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/note="Differential display comparison of untreated and dimethylfumarate treated U937 cells"
17 c 17 g 10 t
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74.18;

 Mismatches

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Pred. No. 26
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EST 30-AUG-2001

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Local Similarity 74.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14 TCCATTCTGCCTGGGTGACAATGCCTT 40
                                                                                                                                                                                                                                                                                                            High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortin/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A1446161 43 bp mRNA linear EST 13-APR-1999
tj079403.x1 NCI_CGAP_Gas4 Homo Sapiens cDNA clone IMAGE:2140852 3/
similar to SW:PRP4_HUMAN P10163 SALIVARY PROLINE-RICH PROTEIN PO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 TCCAAGATGGCTGGGTGGCATTTCCTT 27
                                                                                                                                                                                                                                                                                                                                                           Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                   Trace considered overall poor quality Insert Length: 2385 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                    www-bio.llni.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cgapbs-r@mail.nih.gov
                                                              /note="Organ: stomach; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
                                                                                                                /clone_lib="NCI_CGAP_Gas4"
/tissue_type="poorly differentiated adenocarcinoma with
signet ring cell features"
/lab_host="DH10B"
                       Average insert size 1.69 kb. Life Technologies catalog #: 11549-011"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="PCR was performed on Arabidopsis thaliana lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"
                                                                                                                                                                                                                 /clone="IMAGE:2140852"
                                                                                                                                                                                                                                        /db_xref="taxon:9606
                                                                                                                                                                                                                                                                /organism="Homo sapiens"
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Pred. No. 2e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                  37 CCTTGGTTTCAACACTATCGGAATGTGACTGTCGGGCC 74
                                                                                                                                                                                              HH550482 Incar GSS 13-JU SALK, 071336.42.10.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_071336.42.10.x, DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    42 GGGGACCCCTTTTGCCTGGGGGGCCCGGGGGTTCCTTT 5
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                          BH850482
                                                                                                                                                                                                                                                                                                                                                                        3 CCTTCCTCTGGACTGTAAGAATATGTCTCCAGGGCC 40
                                                                                                                             BH850482.1 GI:21421353
                                                                                                                                                                                                                                                     BH850482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Suzuki, Y.; Yoshitomo-Nakāgawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 GAGCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTT 45
                                                                                 thale cress.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzukiélms.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AU107296 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
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/db_xref="taxon:9606"
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63.2%;
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63.2%;
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Pred. No. 2.
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Pred. No. 2.2e+05;
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PUBMED
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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Unpublished (2001)
                                                                                                                                                    Chen,H., Chrast,R., Rossier,C., Morris,M.A., Lalioti,M.D. and Antonarakis,S.E.
                                                                                                                                                                                                                                               Direct Submission Stylianos E. Antonarakis, Division of Submitted (17-MAR-1995) Stylianos E. Antonarakis, Division of Medical Genetics, University and Cantonal Hospital of Geneva, CMU, 1 rue Michel-Servet, 1211 Geneva, SWITZERLAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chen, H.M., Rossier, C., Chrast, R. and Antonarakis, S.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               H.sapiens DNA for trapped exon (ID HMC37D09), genomic survey
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                           Genome Res. 6 (8), 747-760 (1996)
97011340
                                                                                       exon trapping
                                                                                                                      Cloning of 559 potential exons of genes of human chromosome 21 by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cloning of trapped exons from human chromosome 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note-"PCR was performed on Arabidopsis thaliana lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html" a 7 c 5 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eutheria; Primates; Catarrhini; Hominidae; Homo
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/clone="SALK_071336.42.10.x"
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 Mismatches

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                                                                   24 CTGGGTGACAATGCCTTGGTTTCAACACTATCGGAATGTGA 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
      1 CAGCGATACAGCCACATGAATCCAACACAAACGGCATGGGA 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 45)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BE566921 45 bp mRNA linear EST 15-AUG-
601341047F1 NIH_MGC_53 Homo sapiens cDNA clone IMAGE:3683506 5'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA Library Preparation: CLONTECH Laboratories, Inc. CONA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
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                                                                                                                                                                                                                                                                              //Lisus_type="arcinoma, cell line"
//Lisus_type="arcinoma, cell plage resistant)"
//Lab host="DHIOB (T] phage resistant)"
//note="Organ: bladder; Vector: pDNR-LIB (Clontech);
//note="Organ: bladder; Vector: pDNR-LIB (Clontech);
//note="organ: bladder; Vector: pDNR-LIB (Clontech);
//note="Organ: bladder; Vector: pDNR-LIB (Glontech);
//note="Organ: bladder; Vector: bladder,                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="trapped exon"
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/db_xref="taxon:9606"
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/clone_lib="NIH_MGC_53"
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61.0%;
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Pred. No. 2.
                                                                                                                                                              Score 15.4; DB 10;
Pred. No. 2.6e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17 ATTCTGCCTGGGTGACAATGCCTTGGTTTCAACACTATCGG 57
Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha, Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
                                                                               Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 50)
                                                                                                                                                                                                                                                                                                AU106333 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                    AU106333.1 GI:13555854
                                                                                                                                                                                                                                                                                NBLAN526NR1, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       High quality sequence start: 36 High quality sequence stop: 49.
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Plate: LLCM381 row: k column: 14
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Clone distribution: MGC clone distribution information can be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="Burkitt lymphoma"
/lab_host="DHIOB (phage-resistant)"
/lab_host="DHIOB (phage-resistant)"
/note="Organ: lymph; Vector: poTBS; Site_1: XhoI; Site_2: RocRI; CDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            50 bp mRNA linear EST 30-AUG-200 AUI06615 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone COLF5563, mRNA sequence.
                                                                                                                                                                                                                                                                                                              Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997):
                                                                                                                                                                                                                                                                                                                                                                                                                             Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha, Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Suzuki, r., Yoshitomo-Nakagawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBO Rep. 2 (5), 388-393 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AU106615.1 GI:13556136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Department of Virology Institute of Medical S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 50)
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                                                                                                                       /clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
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/note="Differential display comparison of untreated and
dimethylfumarate treated U37 cells"
11 c 15 g 18 t
                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                       /clone="COLF5363"
                                                                                                                                                                                                                                                                                                      location/Qualifiers
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/db_xref="taxon:9606"
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16.6%;
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61.0%;
Score 15.4; DB 9;
Pred. No. 2.8e+05;
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                        DB 9;
                      Length 50;
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ACCESSION
VERSION
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N57774/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10 GCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAACA 50
                                                                                                                                                                                                                                                                                                                           38 CTTGGTTTCAACACTATCGGAATGTGACTGTCGGGC 73
                                                                                                                                                                                                                                                                             36 CTTGGTTTTATCACCCTCTCTCTCTCTCTATCGCGC 1
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                                                           N57774

39 bp mRNA linear EST 28-JAN-1997
yv56f04.sl Soares fetal liver spleen lNFLS Homo sapiens cDNA clone
IMAGE:246751 3' similar to gbjM87933|HUMAALU364 Human carcinoma
cell-gerived Alu RNA transcript, (rRNA); gb:M96956 EPIDERMAL GROWTH
EAGGNOLITYET DETERM TRANSCRIPT (TRANSCRIPT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The Berkeley Drosophila Genome Project Gene Disruption Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           end
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The P element insertion position is base 29 in the 36 bases. This insertion position refers to the first base of the 8 base target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LSA Building, Berkeley, CA 94720-3200, USA Fax: 5106439947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berkeley Drosophila Genome Project
University of California, Berkeley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Gerald Rubin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Levis,R., Hoskins,R., Liao,G., Mozden,N., Tsang,G., He,Y., Karpen,G., Bellen,H., Rubin,G. and Spradling,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drosophila melanogaster
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N57774.1 GI:1201664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence recovery method was inverse PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fruit fly
                                             FACTOR-LIKE CRIPTO PROTEIN (HUMAN);, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lines Drosophila melanogaster genomic Sequence recovered from 5'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://www.fruitfly.org/about/methods/inverse.pcr.html."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              melanogaster strains each of which contains one or more P\{GT1\} P-element transposon insertion. The resultant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fragment for each strain was directly sequenced to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Inverse PCR was performed on Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                 16.3%;
63.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.2; DB 17;
Pred. No. 2.8e+05;
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  AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11 CACTCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAA 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                          39 CACTCCAGCCTGGGTGACNGAGTGNGACTCTGTCTCAA 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23;
  Clark, M., Johnson, S.L., Lehrach, H., Lee, R., Li, F., Marra, M., Eddy
                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniform; Cyprinidae; Danio.
                                                                                                                                                                                                                                                                        AI601544 46 bp mrNA linear EST 07-JUN-2001 fc08c05.x1 Zebrafish WashU MPING EST Danio rerio cDNA clone IMAGE:3720776 3' similar to SW:EXTN_TOBAC P13983 EXTENSIN PRECURSOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                            Danio rerio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seq primer: m13 -40 forward
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97044478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                     mRNA sequence.
                          (bases 1 to 46)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone-"IMAGE: 246751"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Soares fetal liver spleen 1NFLS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="GDB:3795997"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.2; DB 10
Pred. No. 2.9e+05
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                                                                                Cypriniformes
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COMMENT

TITLE

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ACCESSION
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                                                                                                                                                                                                                                                                                                                                          Match 16.3%; Local Similarity 63.9%;
                                                                                                                                                                                                                    43 GGGGGGGAGAACCCCCTTTTGGGTGGGGCCCAAAG 8
                  A7 bp DNA linear GSS 05-AUG-20 SALK_088536 Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_088536, DNA sequence.

BH862011
                                                                                                                                                                                                                                                                 1 GAGCTGGGAGCACTCCATTCTGCCTGGGTGACAATG 36
BH862011.1 GI:22097337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
Seq primer: T7 ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matthew Clark. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: Genome Systems, St. Louis, Missouri (web address: www.genomesystems.com) (email contact: info@genomesystems.com) and Research Genetics, Huntsville, Alabama (web address: www.resgen.com) (email contact: info@resgen.com) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'S., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood, K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA Library Preparation: Matthew Clark. cDNA Library Arrayed by:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Stephen L. Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and Wilson, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kohn, S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RessourcenZentrumPrimarDatenbank, Berlin, Germany (web address:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WashU
                                                                                                                                                                                                                                                                                                                       Conservative
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314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            zbrafish@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                analysis were selected following oligonucleotide hybridization fingerprinting of arrayed clones from zebrafish late somitogenesis (26 ss), adult liver or embryonic shield stage (5.6 h) libraries. Fingerprint data were used to computationally cluster cDNAs, and a single cDNA from each cluster was chosen for sequencing. In some cases multiple members of the same cluster were sequenced to assess clustering parameters or single clones were sequenced additional times to assess quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COLLCTOT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             digested with Not I and cloned into the Not I and Sal I sites of the pSPORT1 vector (BRL). Library was constructed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       by Matthew Clark (Lehrach lab; ICRF, London and Max Planck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             double-stranded cDNA was ligated to Sal I adaptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Institut fuer Molekulare Genetik, Berlin). cDNAs for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="XL1-blue MRF"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="26 somite embryos, adult livers, shield
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /sex="mixed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:7955"
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    Mismatches

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AZ328471/c
                                                                                                                                                                                                                                                                 REFERENCE
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33 GGAATCAACATTATCGGAAGTAGATTTT 6
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                                       University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                    1 (bases 1 to 48)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,B., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1M0052F16F Mouse 10kb plasmid \tilde{U}UGC1M library Mus musculus genomic clone UUGC1M0052F16 F, DNA sequence.
                                                                                                                                            Unpublished (2000)
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                            Contact: Robert B.
                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                             and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Class: TDNA tagged
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., G., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is single pass sequence recovered from the left border of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Arabidopsis Genome
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A Sequence-Indexed Library of Insertion Mutations in the
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.saik.edu/tdna_protocols.html"
a 10 c 7 g 18 t
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71.4%;
                                                                                                            Weiss
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 CTGCCTGGGTGACAATGCCTTGGTTTCAACACTATC 55
Hillier,L., Lennon,G., Becker,M., Bonaldo,M.F., Chiapelli,B., Chiaspe,S., Dietrich,N., DuBuque,T., Favello,A., Gish,W., Hawkins,M., Hultman,M., Kucaba,T., Lacy,M., Le,M., Le,N., Mardis,E., Moore,B., Morris,M., Parsons,J., Prange,C., Rifkin,L., Rohlfing,T., Schellenberg,K., Soares,M.B., Tan,F., Thierry-Meg,J., Trevaskis,E., Underwood,K., Wohldmann,P., Waterston,R., Wilson,R. and Marra,M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A95D159 mRNA linear EST 01-AUG-199

2003f08:r1 Stratagene colon (#937204) Homo sapiens CDNA clone

IMAGE:566631 5' similar to SW:UCRLHUMAN P31930

UBIQUINOL-CYTOCHROME-C REDUCTASE COMPLEX CORE PROTEIN I PRECURSOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              High quality sequence stop: 48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23;
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Fax: 801 585 7177
                                                                                                                                                                                                                                                Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo 1 (bases 1 to 49)
                                                                                                                                                                                                                                                                                                                                                                                                            EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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Insert Length: 10000 Std Err
                                                                                                                                                                                                                                                                                                                                                                                                                                      AA152159.1 GI:1721211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicallin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      musculus C57BL/6J (male) was obtained from the Jackson
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/clone="UUGC1M0052F16"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /sex="Male
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /strain="C57BL/6J"
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Pred. No. 3.2e+05;
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Local Similarity 64.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        33 TCCANCACCNTCNCCATGGCACTGCCAGGCC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  44 TTCAACACTATCGGAATGTGACTGTCGGGCC 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality possible reversed clone: similarity on wrong strand Insert Length: 1404 Std Error: 0.00 seq primer: -28Mil rev2 from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AI362864 50 bp mRNA linear EST 16-FEB-1999 qy87b11.xl NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2018973 3' similar to gb M19226|MUSRRWB Mouse 28S large subunit rRNA, 5' (rRNA
                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
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Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Wilson RK
www-bio.llni.gov/bbrp/image/image.html
                    Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                      Bonaldo,
                                                                                                                                                                                         Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                               Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AI362864.1 GI:4114485
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                                                                     DNA Sequencing by: Washington University Genome Sequencing Center
                                                                                            cDNA Library Arrayed by: Greg Lennon, Ph.D.
                                                                                                                                             cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mRNA sequence.
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                                                                                                                                                                                                                   cgapbs-r@mail.nih.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cell_line="T84 carcinoma cell line"
/lab_host="SOLR cells (kanamycin resistant)"
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/clone="IMAGE:566631"
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Pred. No. 3.2e+05;
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Trace considered overall poor quality

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 GCCTCGATCAGAAGGACTTGGGCCCCCCACGAGCGG 40
                                                                                                                                                      High quality sequence stop: 40.
                                                                                                                                                                                             Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B
                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (bases 1 to 40)
                                                                                                                                                                                                                                                                                                                                     308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer: -40UP from Gibco quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                              ddunn@genetics.utah.edu
Length: 10000 Std Erro
0339 row: H column: 19
                                                                                                                                                                          plasmid ends
                                                                                                                                                                                                                                                                                                                  USA
                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                    /clone="UUGC1M0339H19"
                                                         /db_xref="taxon:
                                                                            /strain="C57BL/6J"
                                                                                             /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Organ: brain; Vector: pT7T3D-Pac (Pharmacia) with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="NCI_CGAP_Brn25"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:2018973"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Homo sapiens"
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63.9%;
                                                                                                                                   :ion/Qualifiers
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                                                                                                                                                                                                                                                                                                                                               TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE
                                                                                                                                                                                                                                                                     AUTHORS
                                                                                                                                                                                                                                                                                                         JOURNAI
                                                                                                                                                                                                                                                                                                                                                                                                           JOURNAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AUTHORS
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40 CTCTGTTATAGTGTGTTAGTTTTAATCATAAGGAAATGT 2
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                                                                                                        Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion close to or within gene At430820. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German Plant Genomics program designated 'GABI'. Information on line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ", μρ DNA linear GSS 18-JUN-2002 Arabidopsis thaliana T-DNA flanking sequence GK-026F11-013727, genomic survey sequence.
http://www.mpiz-koeln.mpg.de/GABI-Kat/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ROSSO,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat)
                                                                                                                                                                                                                                                                                                                                                      Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut fuer
                                                                                                                                                                                                                                                                                                                                                                                                      Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and Weisshaar, B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Strizhov, N., Li, Y., Rosso, M., Vlehoever, P., Dekker, K., Saedler, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arabidopsis thaliana
                                                                              availability can be found at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for flanking sequence tag based reverse genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Rosso, M., Strizhov, N. and Weisshaar, B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Laboratory Mouse DNA Resource
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Pred. No. 3.5e+05;
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/organism="Arabidopsis thaliana" /strain="Columbia 0"

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA872957 49 bp mRNA linear EST 19-MAY-1998 oh63901.s1 NCI_CGAP_K105 Homo saptens cDNA clone IMAGE:1471728 3 similar to SW:DOPD_HUMAN P30046 D-DOPACHROME TAUTOMERASE;, mRNA
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                                                                                                                                                                                                                                                                                                                                                                                  Insert Length: 400 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                Trace considered overall poor quality Insert Length: 400 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. P.D. CDNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LINL at: www-bio.llni.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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/note-"PCR was performed on DNA from Arabidopsis thaliana
plants (T1) which were transformed with the T-DNA from
vector pAC106. The lines contain one or more T-DNA
lnsertions. The DNA fragment(s) resulting from the PCR
were directly sequenced to determine the genomic sequence
flanking the insertion. Sequences displaying significant
similarity to the A. thalaina nuclear genome sequence were
                                           and Eco RI sites of the modified p7773 vector. Library went through one round of normalization. Library
                                                                                                                                                                                           /clone_lib="NCI_CGAP_Kid5"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                              /clone="IMAGE:1471728"
                                                                                                                                                                                                                                                                                        /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                   RESULT 32
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REFERENCE

AUTHORS

TITLE

JOURNAL

KEYWORDS ACCESSION

ORGANISM

FEATURES

source

RESULT 30 AA872957

BASE COUNT

Query Match

DEFINITION

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Match 16.1%;
Local Similarity 67.7%;
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                                                                                                                                                                                                                                                                                                                                                                                     46 CCATGCTTTGGCTGACACGCTACTGGAACGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                      32 CAATGCCTTGGTTTCAACACTATCGGAATGT 62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jag
Email: ysuzuki@ims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ,H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AU102951 Sugano Homo sapiens cDNA library Homo sapiens cDNA
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                       HUMGS0003674 Human adult (K.Okubo) Homo sapiens cDNA 3', mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBO Rep. 2 (5), 388-393 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                         Homo sapiens
                                                                                                                                         C20586.1 GI:1621696
                                                                                                                                                                                             sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Sugano Homo sapiens cDNA library"
/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ., Morishita,S., Okubo,K., Sakaki
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                                                                                                                                                                                             Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Okubo, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T62015 31 bp mRNA linear EST 14-FEB-1
yb97C06.rl Stratagene lung (#937210) Homo sapiens cDNA clone
IMAGE:79114 5' similar to gb:J03600 ARACHIDONATE 5-LIPOXYGENASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: kousaku@imcb.osaka-u.ac.jp
Human Gene Signature, 3'-directed cDNA sequence. We are not
submitting the same cDNA sequence redundantly to DDBJ since 1993.
For the abundance information of clones with this sequence in this
library and as well as in other 3'-directed libraries, see'
http://www.imcb.osaka-u.ac.jp/bodymap'. The sequences of the clones
Email: est@watson.wustl.edu
                            Tel: 314 286 1800 Fax: 314 286 1810
                                                                      Washington University School of Medicine
                                                                                                                               Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1. (bases 1 to 31)
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Tel: 06-877-5111(ex.3315)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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Contact: Okubo, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   r62015.1 GI:665258
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Similarity 73.1%;
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Pred. No. 3.6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: (
Plate: 0275 row: A column: 10
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 AGCCAGATGGACTCCAGCCTGCCTGG 27
                                                                                                   High quality sequence stop: 36.
                                                                                                                                                                                                                                                                                                        University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
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High qality sequence stops: 18 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Putative
                                                                                                                       Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                               Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                   Unpublished
                                                                                                                                                                                                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                          and Wright, D., Weiss, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ465655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       clone UUGC1M0275A10 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ465655.1 GI:10623696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Insert Length: 489
Seq primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             full length read
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (bases 1 to 36)
                                                                                                                                                                                                                                                                                      308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                  USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
/db_xref="taxon:10090"
                                     /organism="Mus musculus"
                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Stratagene lung (#937210)"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="GDB:4827:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="SOLR cells (kanamycin resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE:79114"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'dev_stage="72 years"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15.9%;
73.1%;
                                                                                                                                                                                                                                                                                                                                                                   (2000)
                                                                                                                                                                                                                                                                                                                                                  Weiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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                                                                                                                                                                              Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GSS 04-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM
                                                                                          source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rocar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  44 TTCAACACTATCGGAATGTGACTGTC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 TTCACAACTGTGGGAAAGTCACTGCC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AA631276 40 bp mRNA linear EST 31-OCT ng92c01.s1 NCI_CGAP_CO9 Homo Sapiens CDNA clone IMAGE:1159716 similar to TR.E239957 E239957 CHROMOSOME XIV READING FRAME ORF
                                                                                                                                                     Insert Length: 639 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
H1gh quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YNL153C. ;, mRNA sequence.
AA631276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            h 15.9%;
Similarity 73.1%;
                                                                                                                                                                                                                                                                                                                                                    CDNA Library Arrayed by: Greg Lennon, Ph.D.

NNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA631276.1 GI:2553887
                                                                                                                                                                                                                                                         Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                        www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissue Procurement: Ilan Kirsch, M.D., Michael R. Emmert-Buck, M.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: M. Bento Soares, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (bases 1 to 40)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gil473214/gb)AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for amplicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="UUGC1M0275A10"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="Male"
/clone="IMAGE:1159776"
                                  /db_xref="taxon:9606"
                                                           /organism="Homo sapiens"
                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hydrodynamically sheared by repeated passage through a 5 inch orifice at constant velocity. The sheared DNA

 Mismatches

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 3.9e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PWD42nv; Purified genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EST 31-0CT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Matches

RESULT 35

ACCESSION VERSION

SOURCE KEYWORDS

REFERENCE

TITLE AUTHORS

JOURNAL

FEATURES

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RESULT 36
AI591257/c
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                                                                                                                                                                                                                                                                                                                                                                                                       FEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMMENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACCESSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    53 ATCGGAATGTGACTGTCGGGCCTCCAGAAAACAT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  #139123/

tt75c06.xl NCI_CGAP_HSC3 Homo sapiens cDNA clone IMAGE:2246602 3/

similar to TR:008808 008808 P140MDIA. ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 43) NCI-CGAP http://www.ncbi.nlm.nih.gov/nciogap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP), National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 15.9%;
Similarity 64.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AI591257.1 GI:4600305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Insert Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA Library Preparation: David B. Krizman, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Procurement: Herbert Morse, M.D., Michael R. Emmert-Buck,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="colon tumor RER+"
/lab_host="DHIOB"
/lab_host="DHIOB"
/note="Organ: colon; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from RER+ colon tumor, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is not normalized. Library was
/note="Organ: bone marrow; Vector: pAMP1; mRNA made from lymphoid tissue, cDNA made by oligo-dT priming. Directionally cloned. Size-selected on agarose gel, average insert size 500 bp. Primary library, non-amplified. cDNA Library Preparation: David B. Krizman, Ph.D. Reference: Krizman et al. (1996) Cancer Research 56:5380-5383."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       constructed by Bento Soares and M. Fatima Bonaldo (Soares4
                                                                                                                                                                                                               /clone="IMAGE:2246602"
/clone_lib="NCI_CGAR_HSC3"
/tissue_type="CD34+, T negative, patient with chronic myelogenous loukemia"
                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="NCI_CGAP_Co9"
                                                                                                                                                                                                                                                                                                                            /db_xref="taxon
                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
                                                                                                                                                                                           'lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 403
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Shinn, P.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        43 CCCCTCTGCATTCATGTCTATGAGTTGCTCTCCCTTCTGTCG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Insert Length: 1116 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   H88764 H9704 H8704 H8704 H8704 H8704 H8704 H8704 H9704 H8704                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              High quality sequence starts: 1
High quality sequence stops: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seq primer: M13RP1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             H88764.1 GI:1071024
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                                                                                                                                                           /lab_host="SOLR cells (kanamycin resistant)"
/lab_host="SOLR cells (kanamycin resistant)"
/note="Organ: ear; Vector: pBluescript SK:; Site_1: EcoRI;
Site_2: XhoI; Reference: Genomics 23, 42-50 (1994) Cloned
unidirectionally. Primer: Oligo dr. Fetal cochiea, normal.
37% of inserts <0.5 kb, 56% 0.5-1.0 kb, 7% >1 kb. Uni-ZAP
XR Vector. Library constructed by N. Robertson, C. Morton.
-5' adaptor sequence: 5' GAATTCGGCACGAG 3' -3' adaptor
sequence: 5' CTGGAGTTTTTTTTTTTTTTTTTTTTT 3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Morton Fetal Cochlea"
/tissue_type="cochlea"
/dev_stage="16-22 week fetus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE: 253103"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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59.5%;
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          Score 14.8; DB 14; Length 47; Pred. No. 4.4e+05;
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                                                                                                                                            8 g
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BH855632
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Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                  1 TCCAAGATGGCTGGTTGGCATTTCCTTTTAATCA 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46 GAGACAGGGTCTNGCCANGTTGCCCAGGTTGGAGTGCAGTGGTT 3
                                                                        EST
                                                                                                                                                                                AU102953 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
                                                                                                 AU102953.1 GI:13552474
                                                                                                                                     AU102953
                                                                                                                                                       ADSE01926, mRNA sequence.
                                                                                                                                                                                                                                   AU102953
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This is single pass sequence recovered from the left border of TDNA. This sequence lies within an annotated exon of At3g05165.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: B58 453 4100 x1752
Fax: B58 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., Gadrinab
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /Clone_lib-"Arabidopsis thaliana TDNA insertion lines"
/note-"PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.saik.edu/tdna_protocols.html"
a 9 c 9 g 20 t
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/db_xref="taxon:37/
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64.78;
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 Mismatches

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Pred. No. 4.5e+05;
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EST 30-AUG-2001

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Gaps

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RESULT 40
H45829
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Suzuki, Y., Yoshitomo-Nakagawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 50)
Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha, Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 H45829 38 bp mRNA linear EST 31-JUL-1995 yo81b05.s1 Soares adult brain N2b4H855Y Homo sapiens cDNA clone IMAGE:184305 3' similar to gb|M87921|HUMALCD120 Human carcinoma cell-derived Alu RNA transcript, (rRNA); gb:X54150_rnal IMMUNOGLOBULIN ALPHA FC RECEPTOR PRECURSOR (HUMAN);, mRNA sequence.
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
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Diverse transcriptional initiation revealed by fine, large-scale
                                                                                                                                                                                                                                                                                                                                                             Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                            , M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R. The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 38)
Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman
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                             Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                   Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
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Insert Length: 1312
                                                                                                                                         High quality sequence starts: 1 High quality sequence stops: 1
                                                                                                                                                                                                                                                                                  Tel: 314 286 1800
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                                                                                                                                                                                                Insert Size: 1312
                                                                                                                                                                                                                       Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                    Fax: 314 286 1810
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/note="plfferential display comparison of untreated and dimethylfumarate treated U937 cells"
18 c 9 g 11 t
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59.5%;
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           Std Error: 0.00
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RESULT 41
AA878693/c
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Location/Qualifiers
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                                                                                          CDNA Library Preparation: M. Bento Soares, Ph.D. CONA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                         Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R. Emmert Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                               Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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  Trace considered overall poor quality Insert Length: 641 Std Error: 0.00
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Insert Length: 641
                                                                          www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA878693.1 GI:2987658
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        provided by Dr. Donald H. Gilden. Tissue The Territory of a 17-18 hours after death which occurred in consequence of a ruptured aortic aneurysm. RNA was prepared from a pool of tissues representing the following areas of the brain: frontal, parietal, temporal and occipital cortex from the left and right hemispheres, subcortical white matter, basal ganglia, thalamus, cerebellum, midbrain, pons and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT773 vector (Pharmacia). Library went through one round of normalization to a Cot = 53. Library constructed by Bento Soares and M.Fatima Bonaldo. The adult brain RNA was provided by Dr. Donald H. Gilden. Tissue was acquired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="GDB:3828657"
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/clone="IMAGE:184305"
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81.0%; Pred. No. 4.
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                                                                                                                                 information can be
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AGE:1492991 3'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  73 CCTCCAGAAAACATTGAGGTG 93
                                                                                                                                                               Possible reversed clone: similarity on w
Seq primer: -28ml3 rev1 ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 40)

Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T. Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylle,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1996)
Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA086682 40 bp mRNA linear EST 15-FEB-1 mn91909.rl Stratagene mouse Tcell 937311 Mus musculus cDNA clone
                                                                                                                                                                                                                                Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                    This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                             Washington University School of MedicineP
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mRNA sequence.
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                                                                                                                                                                                                                                                                              IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA08668
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            house mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IMAGE: 551488 5' similar to SW: CYB5_RAT P00173 CYTOCHROME B5. [1] ;,
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314 286 1810
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                                                                                                                                                                                                                                                                                                                                mouseest@watson.wustl.edu
/clone_lib="Stratagene mouse Tcell 937311"
/tissue_type="Tcell"
                                                     /clone="IMAGE:551488"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               constructed by Bento Soares and M. Fatima Bonaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified p773 vector. Library went through one round of normalization. Library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Mus musculus"
                                                                         /db_xref="taxon:10090"
                                                                                                                                          Location/Qualifiers
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/tlssue_type="2 pooled tumors (clear cell type)"
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Pred. No. 4.9e+05;
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LOCUS
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AU107179 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone COL10026, mRNA sequence.
AU107179

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AU107179.1 GI:13556700
                                                                                                                                                                                                     58 AATGTGACTGTCGGGCCTCCA 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22 GCCTGGGTGACAATGCCTTGGTTTCAACACTATCGGA 58
                                                                                                                                                                                                                                            17;
                                                                                                                                                                      5 AATGCGATGGTCGGGCCCCCA 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tetrahymena thermophila 
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Chicago
920 E. 58th Street, Chicago, IL 60637, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Turkewitz AP
Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EST from Tetrahymena thermophila, strain CU428.1, growing cells Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Turkewitz, A.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: T3
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773 702 3172
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                    /Clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Yector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
proc. Natl. Acad. Sci USA, 98: 8709-8713."
a 12 c 15 g 10 t
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/strain="CU428.1"
/db_xref="taxon:5911"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                         15.7%; Score 14.6; DB 13; Length 46; 81.0%; Pred. No. 5.2e+05;
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 Mismatches

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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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40 bp mRNA linear EST 17-JUN-1998 om63f09.sl NCI_CGAP_GC4 Homo sapiens cDNA clone IMAGE:1551881 3' similar to SW:HBP_HUMAN Q00341 HIGH DENSITY LIPOPROTEIN BINDING
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
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                                                                          Trace considered overall poor quality Insert Length: 815 Std Error: 0.00
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
High quality sequence stop: 1
                                     cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-GAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AA922477
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                                                                                                                                                                                             www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tumor Gene Index
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/db_xref="taxon:9606"
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Pred. No. 5.4e+05;
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    /Clone_lib="Arabidopsis thaliana T-DNA insertion lines"
/note="PCR was performed on DNA from Arabidopsis thaliana
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FEATURES
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                                                                                                                                                                                                                                                                                                                    Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the Left border of the T-NM. It indicates an insertion close to or within gene At5905970. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project, GABI-Kat is part of the German float Context of the CABI-Kat project, GABI-Kat is part of the GABI-Kat project.
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                                                                                                                                                                                                            Plant Genomics program designated 'GABI'. Information on line availability can be found at: http://www.mpiz-koeln.mpg.de/GABI-Kat/.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B.
A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat)
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/clone_lib="NCI_CGAP_GC4"
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/clone="GK-158H06-013215"
                                    /db_xref="taxon:3702"
                                                                        /strain="Columbia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tissue_type="pooled germ cell tumors"
                                                                                                                /organism="Arabidopsis thaliana"
                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15.5%;
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Pred. No. 5.7e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 9;
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RESULT 47
D20665/c
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                                                                                                                                         AI040713/c
                                                                                                                                                                    RESULT 48
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                                                                                                                                                                                                                             32 TCATCTCTATAGGAAGGTGCTGGGCAGGGATC 1
                                                                                                                                                                                                                                                                        45 TCAACACTATCGGAATGTGACTGTCGGGCCTC 76
                                                           A1040713 48 bp mRNA linear EST 28-AUG-10x26a09.sl Soares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:1657432 3', similar to gb:M21121 T-CELL SPECIFIC RANTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             33 AATGCCTTGGTTTCAACACTATCGGAATGTGA 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     32 AATGAGTTTAGCCAAACACTCTCGAAATGGGA 1
AI040713.1 GI:3279907
                                          PROTEIN PRECURSOR (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Osaka University
3-1 Yamada-oka, Suita, Osaka 565, Japan.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yoshinari,H., Arimoto,J. and Matsubara,K.
Institute for Molecular and Cellular Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Okubo, K., Fukushima, A., Yoshii, J., Niiyama, T., Kojima, Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDNA sequencing
Unpublished (1993)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gene expression of human promyelocytic cell line HL60 before and after induction of differentiation. A new application of 3'directed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 45)
Okubo,K., Fukushima,A., Yoshii,J., Nilyama,T., Kojima,Y., Yoshinari,H., Arimoto,J. and Matsubara,K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HUMGS01641 Human promyelocyte Homo sapiens cDNA clone pm1291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          promyelocyte.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Human promyelocyte"
/note="Female, adult, cell_line = HL60, cell_type =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="pm1291"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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65.6%;
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Pred. No. 6e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24;
                                      University of Utah Genome Center University of Utah University of Utah Research 308, Biomedical Polymers Research 308, Biomedical Polymers Research
                                                                                                                                                                                                                        Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 48)
                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 M 0146 B17 F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0146 B17 F, DNA sequence.
Tel: 801 585 5606
                                                                                                                Contact: Robert B.
                                                                                                                                 Unpublished (2000)
                                                                                                                                                             plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZ84615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AZ846157
                                                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                           and Wright, D., Weiss, R
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                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ846157.1 GI:13016065
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 48)
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                                                                                                                                                                                                                                                                                                                                                                                                   nouse mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumor Gene Index
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م
                      USA
                                      Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /dev_stage="8-9 weeks"
/lab_host="DH10B"
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/clone="IMAGE:1657432"
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Pred. No. 6.2e+05;
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49 bp mRNA linear EST 12-FEB-1997
ms30b04.rl Stratagene mouse skin (#937313) Mus musculus cDNA clone
IMAGE:608431 5' similar to SW:RL29_RAT P25886 60S RIBOSOMAL PROTEIN
            Washu-HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Plate: 0146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Insert Length: 10000 Std E Plate: 0146 row: B column:
                                                                                                                                                                                                                                                   Marra,M., Hiller,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T., Gelsel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B.,
                                                                                                            Unpublished (1996)
Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA168179
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Tel: 314 286 1800
                                                                                                                                                                      The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                  Waterston, R.
                                                                                                                                                                                                                     Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA168179.1 GI:1746384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                            house mouse.
                                                                                                                                                                                                                                                                                                                                             (bases 1 to 49)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (g1147321141gblAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ddunn@genetics.utah.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:10090"
/clone="UUGC2M0146B17"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /strain="C5/B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.4; DB 17; Pred. No. 6.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Std Error: 0.00
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                               AUTHORS
                                                                                                                         source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                 Plate: 3526_1_1_1 row: 1 Class: transposon-tagged
                                                                                                                                                                                                                                                                                                        855 California Ave, Palo Alto, CA 94304, USA
                                                                                                                                                                                                                                                                                                                                                                    Contact: Walbot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BH892037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seq primer: -28ml3 rev1 ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fax: 314 286 1810
                                                                                                                                                                                                          Reverse complemented post-ligation sequence from source sequence.
                                                                                                                                                                                                                         Very probable ligation site of ends cut by single endonuclease
                                                                                                                                                                                                                                                 Email: walbot@stanford.edu
                                                                                                                                                                                                                                                                                                                               Stanford University
                                                                                                                                                                                                                                                                                                                                               Department of Biological Sciences
                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2001
                                                                                                                                                                                                                                                                                                                                                                                                          Maize genomic sequences found using engineered RescueMu transposon
                                                                                                                                                                                                                                                                                                                                                                                                                             Walbot, V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clade; Panicoideae; Andropogoneae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BH892037.1 GI:22223095
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     uea mays.
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                                                                                                                                                                                                                                                                   650 723 2227
650 725 8221
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="whole skin"
/dev_stage="11 weeks old"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /sex="females
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Stratagene mouse skin (#937313)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /strain="C57BL
/dev_stage="adult"
                     /tissue_type="leaf"
                                         /db_xref="taxon:4577"
/clone_lib="3526 - RescueMu Grid K"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="IMAGE:608431"
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                                                                                /cultivar="mixed background W23/A188/B73"
                                                                                                      /organism="Zea mays"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Mus musculus"
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                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15.5%; Score 14.4; DB 9; Length 49; 60.0%; Pred. No. 6.3e+05;
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34 CCAGCCCTCGGGTTCAGGAATCTCGGAAAGCG 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 50)

Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T. Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Marra M/Mouse EST Project
WashU-HHMI Mouse EST Project
WashIngton University School of Medicinep
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA674026 Incar EST 26-NOV-1
vp70d06.rl Knowles Solter mouse blastocyst By Mus musculus cDNA
clone IMAGE:1082027 5' similar to gb:X13661 Mouse mRNA for
elongation factor 1-alpha (MOUSE); mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                       Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
TMACE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA674026.1 GI:2647308
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314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          units. For more information on RéscueMu, go to the web site 'www.zmdb.iastate.edu' and follow the links for 'RescueMu.' Grid K was grown at Molokai, Hawaii in Winter 2000-2001. DNA was extracted from leaf punches, double digested using BamHI and BglII, and ligated to form circular plasmids. DH10B cells were transformed and then screened on LB plates with ampicillin."
                           designed to allow plasmid rescue from total genomic DNA. Mu elements insert preferentially into transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="organ: leaf; Vector: RescueMu (engineered from
pBlueScript backbone); Site_1: BamHI; Site_2: BglII;
RescueMu is a 4.9 kb, modified maize Mu transposon
cloned into the NotI/SalI sites of a pSPORT vector (Life
                                                                                                                    /clone_lib="Knowles Solter mouse blastocyst B3"
/tissue_type="blastocyst"
/dev_stage="embryo (pre-implantation)"
/lab_host="pH10B"
                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J x DBA/2J F1"
                                                                                                                                                                                                                              /clone="IMAGE:1082027"
                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /lab_host="DH10B"
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                                                                                                                                                                             DEFINITION
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                                                                                                       VERSION
                                                                                                                               ACCESSION
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                                                                                                                                                                                                                                                       RESULT 54
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                         11 CACTCCATTCTGCCTGGGTGACAA 34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
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                                                                                                                                                               AU103596 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone CAE03627, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,r., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
,S. Construction and characterization of a full length-enriched and
a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                 Homo sapiens
                                                                                                                                               AU103596
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
EMBO Rep. 2 (5), 388-393 (2001)
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50 bp mRNA linear EST 30-AUG-200
AU102972 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
COL08352, mRNA sequence.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                          human.
                                                                                                                 AU103596.1 GI:13553117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (bases 1 to 50)
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/note="Differential display comparison of untreated and
dimethylfumarate treated U937 cells"
a 12 c 16 g 14 t
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/db_xref="taxon:9606"
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 Mismatches

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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                        Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzukiéims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
                                                                                                                                                                                                                                                                                                                                                                                                                                      Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites EMBO Rep. 2 (5), 388-393 (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COLF6124, mRNA sequence
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Similarity 60.0%;
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                                                                                                                                                                                                                                                                                                                                                                                          Contact: Yutaka Suzuki
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5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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/clone_lib="Sugano Homo sapiens cDNA library" /note="Differential display comparison of untreated and dimethylfumarate treated U937 cells"
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/clone=lb="Sugano Homo sapiens cDNA library"
/clone=plferential display comparison of untreated and dimethy. Itumarate treated U937 cells"
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/db_xref="taxon:9606"
                                                                                /clone="COLF6124"
                                                                                                        /db_xref="taxon:9606"
                                                                                                                              /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                         18 TTCTGCCTGGGTGACA 33
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                                                                                                                                                                                                                                                                                                                                                                                            49 TTCAGCCTGGGTGACA 34
                                                                                                                                                                             AV836264 K. Sato unpublished cDNA library: Hordeum vulgare subsp. vulgare seedling leaves second leaf stage Hordeum vulgare subsp. vulgare cDNA clone basd2001, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Suzuki, Y., Yoshitomo-Nakagawa, K., Maruyama, K., Suyama, A. and Sugano, S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AU106860
50 bp mRNA linear EST 30-AUG-200 AU106860 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone COL03340, mRNA sequence.
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                        Hordeum vulgare subsp. vulgare.
Hordeum vulgare subsp. vulgare
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
                                                                                                        EST
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H., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki
Y., Nakamura,Y., Suyama,A. and Sugano,S.
Diverse transcriptional initiation revealed by fine, large-scale
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooideae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzukieims.u-tokyo.ac.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBO Rep. 2 (5), 388-393 (2001)
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1. (bases 1 to 50)
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/note="bifferential display comparison of untreated and dimethylfumarate treated U937 cells"
10 c 11 g 22 t
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/db_xref="taxon:9606"
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93.8%;
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Pred. No. 6.3e+05;
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WashU-NCI human EST Project
Unpublished (1997)
                                                                                                                                 This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
Seq primer: -40Up from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AI582089 31 bp mrNA linear EST 06-APR-1999 ar9609.x1 Barstead colon HPLRB7 Homo sapiens cDNA clone IMAGE:2173456 3' similar to TR:021792 021792 CYTOCHROME B ;contains element MSR1 repetitive element ; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                           Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                              Contact: Wilson RK
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                                                                                                                      High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                Tel: 314 286 1800
Fax: 314 286 1810
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Okayama University, Barley Germplasm Center
Chuo 2-20-1, Kurashiki, Okayama 710-0046, Japan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: kazsato@rib.okayama-u.ac.jp,
URL:http://www.rib.okayama-u.ac.jp/barley/
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/clone="IMAGE:2173456"
                        /db_xref="taxon:9606
                                           /organism="Homo sapiens"
                                                                                                 Location/Qualifiers
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/clone="basd2o01"
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                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0085 row: M column: 1:
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                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 32.
                                                                                                                                                                                                                                                                                                                                                             Seq primer: CGTTGTAAAACGACGGCCAGTClass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus (bases 1 to 32)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2M0085M12F Mouse 10kb plasmid UUGC1M library Mus musculus genomic cione UUGC2M0085M12 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 TGGGAGCACTCCATTCTGCCTGGGTGA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ816444.1 GI:12986352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 70.4
19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         308,
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801 585 7177
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20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                            musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Laboratory Mouse DNA Resource
                                                                                                                                                                                                          /db_xref="taxon:10090"
/clone="UUGC2M0085M12"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                             /note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                             /sex="Male"
                                                                                                                                                                             /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                     /strain="C57BL/6J"
                                                                                                                                                                                                                                                                     /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Barstead."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3']; double-stranded cDNA was ligated to Eco RI adaptors [5' AATTCACTAGTAAT 3' and 5' ATTACTAGTG 3'], digested with Not I and cloned into the Not I and Eco RI sites of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the modified pT7T3 vector. Library constructed by Bob
                                                                                                                                                                                                                                                                                                                     .ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local Similarity
                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 TTCTGCCTGGGTGACAATGCCTTGGTT 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               36 bp DNA linear GSS 02-MAY-200 SALK_065834 Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_065834, DNA sequence.
BH814159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 TTGTTCCTGGGGGATAATTGCTTTGAT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BH814159.1 GI:20393445
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  thale cress.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This is single pass sequence recovered from the left border of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Prednis, L., Shinn, P. Zimmerman, J. and Ecker, J.R. Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (bases 1 to 36)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarcose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 (g114732114|gb1AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                              13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ecker@salk.edu
                                                                                                                                                                                                                    /clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
                                                                                                     be found at http://signal.salk.edu/tdna_protocols.html*
4 c 8 g 11 t
                                                                                                                                                                   directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
                                                                                                                                                                                                                                                                                                                                            /clone-"SALK_065834"
                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:3702"
                                                                                                                                                                                                                                                                                                                                                                                                   /strain-"Columbia
                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Arabidopsis thaliana"
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15.3%;
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Score 14.2; DB 17 Pred. No. 6.5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.2; DB 17;
Pred. No. 6.1e+05;
                           DB 17;
                           Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 32;
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SOURCE

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                                                                                ORIGIN
                                                                                                              BASE COUNT
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High quality sequence stop: 37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 37)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
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Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84112
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0276 row: O column: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nouse mouse
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                                                                                                        10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gli4732114gb)AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                      was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 musculus C57BL/6J (female) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="Female"
/lab_host="E. coli strain XL10-Gold, T1-resistant, F-"
/lab_host="E. coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC2M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:10090"
/clone="UUGC2M0276023"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
15.3%; Score 14.2; DB 17; 62.9%; Pred. No. 6.5e+05;
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 Mismatches

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                                Length 37;
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Other_ESTs: fd23a09.y1
Other_total Stephen L. Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    54 TCGGAATGTGACTGTCGGGCCTCCAGAAAACATTG 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA Library Preparation: Matthew Clark. CDNA Library Arrayed by: Matthew Clark. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: Genome Systems, St. Louis, Missouri (web address: www.genomesystems.com) (email contact: info@genomesystems.com) and Research Genetics, Huntsville, Alabama (web address: www.resgen.com) (email contact: info@genomesystems.com) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Stephen L. Johnson
Washington University School of Medicine
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Clark,M., Johnson,S.L., Lehrach,H., Lee,R., Li,F., Marra,M., I, S., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Unde,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40 bp mRNA linear EST 25-AUG-1999 fd23a09.xl Zebrafish WashU MPIMG EST Danio rerio cDNA 3' similar to SW: F934 HUMAN Q13347 EUKARYOTIC TRANSLATION INITIATION FACTOR 3 A1974168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: T7 ET from Amersham
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RessourcenZentrumPrimarDatenbank, Berlin, Germany (web address:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WashU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and Wilson, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Accinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Danio rerio
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314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zebrafish EST Project 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           zbrafish@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R.
analysis were selected following oligonucleotide hybridization fingerprinting of arrayed clones from zebrafish late somitogenesis (26 ss), adult liver or embryonic shield stage (5.6 h) libraries. Fingerprint data were used to computationally cluster cDNAS, and a single cDNA from each cluster was chosen for sequencing. In some cases multiple members of the same cluster were sequenced to assess clustering parameters or single clones
                                                                                                                                                                                                               double-stranded cDNA was ligated to Sal I adaptors (BRL), digested with Not I and Sal I sites of the pSporTl vector (BRL). Library was constructed by Matthew Clark (Lehrach lab; ICRF, London and Max Planck Institut fuer Molekulare Genetik, Berlin). cDNAs for EST
                                                                                                                                                                                                                                                                                                                                                 /tissue_type="26 somite embryos, adult livers, shield
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Zebrafish WashU MPIMG EST"
                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="XL1-blue MRF"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="mixed"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Danio rerio"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ion/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Insert Length: 3181 Std Erro
Seq primer: m13 -40 forward
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence starts: 1
High quality sequence stops: 1
Source: IMAGE Consortium, LLNL
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yx06e05.sl Soares melanocyte 2NbHM Homo sapins CDNA clone
IMAGE:260960 3' similar to gbh M87915 HUMALNE54 Human carcinoma
cell-derived alu RNA transcript, (rRNA); gb:L06133
COPPER-TRANSPORTING ATPASE 1 (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ,R., Williamson,A., Wohldmar
The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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314 286 1810
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                                              double-stranded cDNA was size selected, ligated to Eco'RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT/73 vector (Pharmacia). Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from normal foreskin melanocytes (FS374) was kindly provided by T. Anthony P. Albino." a 12 c 13 g 9 t 1 others
                                                                                                                                                                                                                             /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="GDB:3870602"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Soares melanocyte 2NbHM"
                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:260960"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
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70.4%;
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 Mismatches

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COMMENT

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     BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ663145/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 AGCTGGGAGCACTCCATTCTGCCTGGG 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1M0542D09R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0542D09 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ663145.1 GI:11800291
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Similarity 70.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lass: plasmid ends
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     13 a
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Length: 10000 Std Erro
0542 row: D column: 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114)gbhAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide Kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                                                                                                                                                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                          adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /sex="Male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone-"UUGC1M0542D09"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"

 Mismatches

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                                                                                                                                                                                                                                                                RESULT 66
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                                                                                       KEYWORDS
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DEFINITION
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Best Local 9
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Local Similarity 62.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  38 TATCAGCATGTAGCCCTGTAGCCTTCACAACATAT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   52 TATCGGAATGTGACTGTCGGGCCTCCAGAAAACAT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                           21 TGCCTGGGTGACAATGCCTTGGTTTCA 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to give a tight size distribution (
4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects.
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TRED927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 ISA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                            5 TGCCTCAGTTACAATATCTTCTTTCCA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  petails of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genomic survey sequence
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                                                                                Zea mays
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AL476863.1 GI:11843427
                                                                                                                                                          вн890613.1 GI:22221585
                                                                                                                                                                                        вн89061
                                                                                                                                                                                                               DNA sequence.
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                                                                                                           Zea mays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: nelsayed@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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/strain="TREU927"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:5691"
/clone="203e07"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.2; DB 17
Pred. No. 6.8e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             36 GGACTCCAGCCTCGGGTTCAGGAATCTCGGAAAG 2
Contact: Kazuhiro Sato
Research Institute for Bioresources
Research Institute for Bioresources
Okayama University, Barley Germplasm Center
Chuo 2-20-1, Kurashiki, Okayama 710-0046, Japan
Email: kazsatoerib.okayama-u.ac.jp,
URL:http://www.rib.okayama-u.ac.jp/Darley/
Sato,K., Saisho,D., Takeda,K., Shini,T. and Kohr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27 GGTGACAATGCCTTGGTTTCAACACTATCGGAATG 61
                                                                                                                                                                                                                                                                                                                                                   Hordeum vulgare subsp. vulgare. Hordeum vulgare subsp. vulgare
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AV833659 K. Sato unpublished cDNA library: Hordeum vulgare subsp. vulgare shoots germination Hordeum vulgare subsp. vulgare cDNA
                                                                                                                                                                                   Unpublished (2001)
                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Štreptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Lillopsida; Poales; Poaceae; Pooideae
                                                                                                                                                                                                   Barley EST sequencing project in NIG and Okayama Univ
                                                                                                                                                                                                                                                                                                                                                                                                                                       AV833659.1 GI:14525748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  clone bags1h21, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        855 California Ave, Palo Alto, CA 94304, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Department of Biological Sciences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2001 Contact: Walbot V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Maize genomic sequences found using engineered RescueMu transposon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 45) Walbot, V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Reverse complemented post ligation sequence from source sequence. Plate: 3526_1_14_1 row: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: walbot@stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stanford University
                                                                                                                                                                                                                                                                                    Triticeae;
                                                                                                                                                                                                                                                               (bases 1 to 46)
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650 725 8221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 units. For more information on Rescuedu, go to the web site 'www.zmdb.iastate.edu' and follow the links for 'Rescuedu.' Grid K was grown at Molokai, Hawaii in Winter. 2000-2001. DNA was extracted from leaf punches, double digested using BamHI and BglII, and ligated to form circular plasmids. DH10B cells were transformed and then screened on LB plates with ampicillin."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site_1: BamHI; Site_2: BglII; RescueMu is a 4.9 kb, modified maize Mu transposon designed to allow plasmid rescue from total genomic DNA. Mu elements insert preferentially into transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /cultivar="mixed background W23/A188/B73"
/db_xref="taxon:4577"
/clone_lib="3526 - RescueMu Grid K"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="leaf"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Zea mays"
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                                                                                                                                                                                                                                                                                    Hordeum.
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     and Kohara, Y. Direct
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AV965348
                                            AU104251
                                                             RESULT 69
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                                                                                                                                              40 TGGTTTCAACACTATCGGAATGTGACTGTCGGGCCTCCAGAAA 82
AU104251 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
                                                                                                           AU104251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AV965348 47 bp mRNA linear EST 14-MAR-2 AV965348 Nori Satoh unpublished cDNA library, young adult Ciona intestinalis cDNA clone ciad19p03 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: satoh@ascidian.zool.kyoto-u.ac.jp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 81-75-753-408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Nori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T. Expressed genes in Ciona intestinalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 GTTTTGAAGAACCCATTCTACGAGNATGGAAATGCC 37
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/tissue_type="shoots"
                                                                                                                                                                                                                                                                                                         /dev_stage="young adult"
                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Nori Satoh unpublished cDNA library, young
                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:7719"
                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Ciona intestinalis"
                                                                                                                                                                                                                                                                                                                            /tissue_type="whole animal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
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/db_xref="taxon:112509"
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    Mismatches

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Pred. No. 7.2e+05;
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                     EST 30-AUG-2001
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           Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki.y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2), 149-156 (1997).
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                                                                                                                                                                                                                                                    1 (bases 1 to 50)
Suzuki, Y., Taira, H., Tsunoda, T., Mizushima-Sugano, J., Sese, J., Ha
H., Ota, T., Isogai, T., Tanaka, T., Morishita, S., Okubo, K., Sakaki
Y., Nakamura, Y., Suyama, A. and Sugano, S.
Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
mapping of mRNA start sites
EMBO. Rep. 2 (5), 388-393 (2001)
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Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Department
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                          Contact: Yutaka Suzuki
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
S. Construction and characterization of a full length-enriched and
a 5'-end-enriched CDNA library. Gene 200 (1-2), 149-156 (1997).
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Diverse transcriptional initiation revealed by fine, large-Scale mapping of mRNA start sites
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AZ443784
                  1M0238G02F Mouse 10kb plasmid ŪUGC1M library Mus musculus genomic clone UUGC1M0238G02 F, DNA sequence.
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Similarity 58.1%;
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dimethylfumarate treated U97 cells"
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Pred. No. 7.5e+05;
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Pred. No. 7.5e+05;
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                                                                      DNA
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                                                                                                                                                                                                                     4 CCATCCTGACTTTGTGACATTG 25
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              BM396571
                                                  BM396571 29 bp mRNA linear EST 17 5009-0-22-D05.t.1 Chilcoat/Turkewitz cDNA (large fraction)
                                 Tetrahymena thermophila cDNA, mRNA sequence.
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Insert Length: 10000 Std Error:
Plate: 0238 row: G column: 02
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B.
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Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 26)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNO42 (gil4732114(gb)RF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll X110-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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77.3%;
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Pred. No. 6.6e+05;
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                                                                      Email: ddunn@genetics:utah.edu
Insert Length: 10000 Std Erro
Plate: 0311 row: 0 column: 2
                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                           University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1M0311024F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0311024 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 AGCTGGGAGCNCTCC 20
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High quality sequence stop: 35
                         Class: plasmid ends
                                                     Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                    and Wright, D., Weiss, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 773 702 4374 Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2002)
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                                                                                                                                                                                                                                                                                                                  Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
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1 (bases 1 to 35)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               University of Chicago
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  house mouse.
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/note="Vector: BlueScript2 SK; betails on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
8 c 11 g 3 t 1 others
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/strain="CU428 1"
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                                                                                                                                                                                                                                                                                                                  Weiss

    Mismatches

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Pred. No. 6.9e+05;
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1; Indels Length 29;

0; Gaps

GSS 05-OCT-2000

Std Error: 0.00

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DR24J19S/c
LOCUS
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                                                                                                                                                                                                                                                                                        Further details: http://www.sanger.ac.uk/Projects/D_rerio/
                                                                                                                                                                                                                                                                                                                                        Campus, Hinxton, Cambridgeshire, CB10 1SA, UK. E-mail contact: humquery@sanger.ac.uk Unpublished
This sequence was generated from the SP6 end of BAC 24J19. 24J19 : part of the Daniokey Pilot BAC Library created by R. Plasterk and
                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (06-JUN-2002) The Sanger Institute, Wellcome Trust Genome Campus, Hinxton, Cambridgeshire, CB10 1SA, UK. E-mail contact:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Humphray, S.J., Huckle, E. and Hunt, S.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes; Cyprinidae; Danio.

1 (bases 1 to 35)
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a
                                                                      /tissue_type="Testis"
/note="vector pIndigoBAC-536"
6 c 9 g 8 t
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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                                                                                                                                                             /clone="DKey-24J19"
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                                                                                                                                                                                                          /organism="Danio rerio"
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  15.1%; Score 14; DB 17;
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  15.1%; Score 14; DB 17; Length 36;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Etd Error: 0.00
Plate: 0089 row: C column: 09
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 36)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     University of Utah Genome Center University of Utah
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2M0089C09R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss
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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114 [gb]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampiciliin resistance."
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/clone="UUGC2M0089C09"
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High quality sequence stop: 37.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0029 row: M column: 20
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (g114732114/gblAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="UUGC2M0029M20"
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        15.1%; Score 14; DB 17; Length 37;
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0024h05.x1 Soares_NSF_F8_9W_OT_PA_F1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: ggapbs-r@mail.nin.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 TCTGGTTGGTTGACACTGTTGTTCTTCCAA 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                High quality sequence stop: 1.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Insert Length: 826
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AI088003.1 GI:3427036
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                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         libraries were mixed, and ss circles were made in vitro.
Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 5 libraries. The pools consisted of the following libraries and cloneiDs: Soares NbHSF pool 1:
309384-310919, 323208-335895 Soares NbHF pool 1:
145032-147335, 147720-148103, 148872-149255, 15002-150407, 151176-152327 Soares NbHFR-pool 1:
758280-760583, 772104-774407 Soares NbHPA pool 1:
758280-760583, 772104-774407 Soares NbHPA pool 1:
304776-306311, 320136-322823, 326280-32663 Soares NbHOT pool 1: 723720-726407, 739080-740999 Subtraction by Bento Soares and M. Fatima Bonaldo."
7 a 9 c 11 g 11 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:1567161"
/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
/lab_host="DH10B"
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39 bp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0339 row: H column: 19
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases I to 40)
1 (bases I to 40)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1M0339H19R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0339H19 R, DNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Nori Satoh
Department of Zoology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T. Expressed genes in Ciona intestinalis Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ciona intestinalis
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Class: plasmid ends
                                                                                                                                                                                               University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                          Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kyoto University
Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EST
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                                                                                                                                                                                                                                                                                          plasmid inserts
                                                                                                                                                                                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                   and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ501159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: satoh@ascidian.zool.kyoto-u.ac.jp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Phlebobranchia; Cionidae; Ciona.
1 (bases 1 to 39)
                                                                                                           Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZ501159.1 GI:10682475
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/clone_lib="Nori Satoh unpublished cDNA library, egg"
/tlssue_type="whole animal"
                                                                                                                                                                            Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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8 c 9 g 10 t
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/db_xref="taxon:7719"
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                                                                                                                                                                                                                                                                                                                                                                                        TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly, Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GACATTTCCTTATGATTAAAACTAACACACTATAACAG 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 40.
Class: plasmid ends
                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0245 row: G column: 17
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                              University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 43)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1M0245G17F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0245G17 F, DNA sequence.
                                                                                                                                                                                                                                                                                                         Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                           plasmid inserts
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Similarity 60.5%;
                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                  84112
                                                                                                                                                                                                                                                                                                                                                                                                                and Wright, D., Weiss, R.
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/clone="UUGC1M0339H19"
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                                                                                                                                                                                                                                                                                                            Weiss
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                                                                                     Std Error: 0.00
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                                                                                                                             Sequencing primer: SP6
This sequence is from a Xenopus Gene Collection (XGC) library constructed by Aaron M. Zorn.
                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 44)
Huckle, E., Taylor, R., Ashurst, J.L., Zorn, A.M. and Rogers, J.
Sanger Xenopus tropicalis EST project 2001 (10_2001)
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                         Contact: Huckle E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9 AGCACTCCATTCTGCCTGGGTGACAATGCC 38
                                                                                                                                                                                                           Sanger Xenopus tropicalis EST project 2001
TROPICALIS_SEQUENCE_ID: TGas012m13.sp6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Silurana tropicalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNA sequence
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Location/Qualifiers
                                                                                                                                                                                                                                                                                            Hinxton,
                                                                                                                                                                                                                                                                                                                    Sanger Centre
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Xenopodinae; Silurana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    western clawed frog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AL629278.1 GI:16598761
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                                                                                                                                                                                                                                                                trop@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwnA2 (gil4732114|gb|AFT29072-1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XLLO-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="XGC-gastrula"
                          /clone="TGas012m13"
                                            /db_xref="taxon:8364"
                                                                 /organism="Silurana tropicalis"
                                                                                                                     Location/Qualifiers
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66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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Pred. No. 8.2e+05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44 bp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA
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AZ779267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62 TGACTGTCGGGCCTCCAGAAAACATTGAGG 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 46.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0015 row: O column: 21
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ,M., Rose,M., Rose,R. and Wright,D.,Weiss,R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 TGTCTGGCGGGTCTTGGCGAAACACTGGGG 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 46)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ779267.1 GI:12909749
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative
                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                       musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                          /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/clone="UUGC2M0015021"
                                                                                                                                                                                                                                                                                                                                                                                                                                                   /strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /dev_stage="gastrula (stages 10.5-13 mixed)"
/lab_host="Escherichia coli XLI-blue"
/lab_host="Escherichia coli XLI-blue"
/note="Vector: pCS107; Site_1: EcoRI; Site_2: NotI; cDNA
was oligo dT primed from 5ug of poly A+ RNA from stages
10-13 gastrulae. EcoRI-NotI cut cDNA was then ligated
cod "ROS107 with EcoRI at the 5' end and NotI at the 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Mus musculus"
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66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No.
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AZ785009

SOURCE

KEYWORDS VERSION

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BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah Genome Center 
University of Utah
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Similarity 60.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Length: 10000 Std Errc
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative
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                                                                                                                                                                                                                                                          musculus C57BL/6J (male) was obtained Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:10090"
/clone="UUGC2M0028B05"
                                                                                                                                                                                                                                                                                                                 /note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                               /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                              /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Mus musculus"
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suzuki,Y., Taira,H., Tsunoda,T., Mizushima-Sugano,J., Sese,J., Ha, Ha, G., Ota,T., Isogai,T., Tanaka,T., Morishita,S., Okubo,K., Sakaki,Y., Nakamura,Y., Suyama,A. and Sugano,S. Diverse transcriptional initiation revealed by fine, large-scale mapping of mRNA start sites.
AU105649
         50 bp mRNA linear EST 30-AUG-200 AUJ05649 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone HSIO2850, mRNA sequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jag
Email: ysuzuki@ims.u-tokyo.ac.jp
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/note="bifferential display comparison of untreated and
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/clone | 1 + - " =
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Diverse transcriptional initiation revealed by fine, large-scale
                                                                                                                                                                   Email: cliew@rics.bwh.harvard.edu
Seq primer: GAAATTAACCCTCACTAAAGGG.
                                                                                                                                                                                                                                                                                       Brigham and Women's Hospital
Harvard Medical School
                                                                                                                                                                                                                                                                                                                                                                Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                    cDNAs from fetal heart (1996)
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
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                                                                                                                                                                                                                                                                75 Francis St. Boston, MA 02115, USA
                                                                                                                                                                                                                                                                                                                                              Contact: Liew CC
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/clone_lib="Sugano Homo sapiens cDNA library"
/clote="Differential display comparison of untreated and dimethylfumarate treated U937 cells"
a 13 c 21 g 6 t
/clone_lib="Human fetal heart, Lambda ZAP Express"
/lab_host="E. coli XL1-Blue"
                                                   /clone-"J2084"
                                                                        /db_xref="taxon:9606"
                                                                                               ∕organism≃"Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              госат
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17 ATTCTGCCTGGGTGACAATGCCTTGGTTTC 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 50.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1M0369A14F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0369 row: A column: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 50)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ580938.1 GI:11695451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clone UUGC1M0369A14 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nouse mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       predigested lambda ZAP Express."
a 17 c 10 g 13 t
                                                                                                                                                                                                                                                                                                             /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   weeks). cDNA was synthesized using a XhoI-Oligo dT adaptor-primer. EcoRI adaptors were ligated, follodigestion with XhoI, for directional cloning into
                                                                                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                  /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                               /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: Lambda ZAP Express; Site_1: EcoRI; Site_2:
XhoI; mRNA was purified from human fetal hearts (8-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="UUGC1M0369A14"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /strain="C57BL/6J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Std Error: 0.00
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inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to

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BASE COUNT
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                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JOURNAL
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                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             48 ACACTATCGGAATGTGACTGTC 69
                                            29 TGACAATGCCTTGGTTTCAACACTA 53
7 TTACAATGCTTTTGTAACACCATTA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BH789431.1 GI:19882529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BH789431 31 bp DNA linear GSS SALK_029414.47.75.x Arabidopsis thaliana TDNA insertion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This is single pass sequence recovered from the left border of TDNA. This sequence lies within 300 bases of the 3' end of At1g17870.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Arabidopsis thaliana genomic clone SALK_029414.47.75.x, DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1 (bases 1 to 31)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ecker@salk.edu
                                                                                                                                                                                                                                                               each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

12 c 10 g 18 t
                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
                                                                                                                                                                                                                                                                                                                                                                                                             /strain="Columbia 0"
/db_xref="taxon:3702"
/clone="SALK_029414.47.75.x"
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77.3%;
                                                                                                                   14.8%; Score 13.8; DB 17; 72.0%; Pred. No. 8.4e+05;
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0; Mismatches 7;
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RESULT 90
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                                                                                                                                                                                                          WashIngton University School of MedicineP
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculius

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Description of the control of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Danio rerio
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Email: mouseest@watson.wustl.edu This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                Unpublished (1996)
Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Marra,M., Hillier,L., Allen,M., Bowles,M., Districh,N., Dbuque,T., Gelsel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Campus, Hinxton, Cambridgeshire, Chumquery@sanger.ac.uk Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (06-JUN-2002) The Sanger Institute, Wellcome Trust Genome Campus, Hinxton, Cambridgeshire, CB10 1SA, UK. E-mail contact:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cypriniformes;
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                                                                                                                                     Tel: 314 286 1800 Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                            The WashU-HHMI Mouse EST Project
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DR11H23T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA048244.1 GI:1527914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             details: http://www.sanger.ac.uk/Projects/D_rerio/
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/tissue_type="Testis"
/note="vector pIndigoBAC-536"
12 c 7 g 7 t
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72.0%;
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Pred. No. 8.6e+05;
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FEATURES
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                         Class: plasmid ends
                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0036 row: M column: 13
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                 Contact: Robert B
                                                                                                                                                                                                                                                                                                      plasmid inserts
Unpublished (20
                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                        Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 39)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GAGCTGGGAGCACTCCATTCTGCCTGGGTGACA 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Trace considered overall poor quality Possible reversed clone: similarity on wrong strand Seq primer: -28M13 rev2 from Amersham
High quality sequence stop: 39.
                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone UUGC1M0036M13 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1M0036M13R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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                                                                                                                                                                                          USA
                                                                                                                                                                   585 5606
                                                                                                                                                                                                          Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT/T3 vector. Library went through one round of normalization, and was constructed by Bento Soares and M.Fatima Bonaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="embryo"
/dev_stage="13.5-14.5dpc total fetus"
/lab_host="DHIOB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T 3'], on equal amounts of mRNA from 2 13.5dpc and 2 14.5dpc embryos [total RNA provided by Minoru Ko, Wayne State Univ., from 2 ]; double-stranded cDNA was ligated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Soares mouse embryo NbME13.5 14.5"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:477294"
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                                                                                              Std Error: 0.00
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seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 1.
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                                                                                                                  Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHAIN PRECURSOR. ;, mRNA sequence
                                                                                                                                                                                                                         www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA642607.1 GI:2567825
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance. ^{\rm m}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
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72.0%;
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                                                                                                                                                                                                                                               High quality sequence stop: 40.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                Email: ddunn@genetlcs.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0545 row: L column: 22
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 40)
                                                                                                                                                                                                                                                                                               Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ665156.1 GI:11802302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ665156 40 bp DNA linear GSS 14-DEC-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           308,
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                   /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                         /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
  Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Organ: prostate; Vector: pT7T3D-Pac (Pharmacia)
with a modified polylinker; 1st strand cDNA was prepared
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/db_xref="taxon:9606"
/clone="IMAGE:1158011"
                                                                /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                             /sex="Male"
                                                                                                           /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                   /clone-"UUGC1M0545L22"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from normal prostate bulk tissue, and was then primed with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="NCI_CGAP_Pr22"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="normal prostate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14.8%;
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                                                                                                                                                                                                                                                                                                                                                             Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA sequence.
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                                                                                           ORIGIN
                                                                                                                BASE COUNT
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                   Best Local Similarity
                                           Query Match
Matches
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Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TRE0927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 TITATACCCTTTTGAAATGTGGCTG 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TA381G12Q 41 bp DNA linear GSS 13-DITA STUCE Sheared genomic DNA clone 381g12, reverse sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Trypanosoma brucei.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AL497699.1 GI:11873421
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   give a tight size distribution ( k). The k + k i method used for the library construction is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases
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                                                                                                                                                                                               /db_xref="taxon:5691"
/clone="381g12"
                                                                                                                                                                                                                                                                        /strain="TREU927"
                                                                                                                                                                                                                                                                                                        /organism="Trypanosoma brucei"
                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                            14.8%; Score 13.8; DB 17
88.2%; Pred. No. 9.5e+05;
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Pred. No. 9.4e+05;
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                                                             DB 17;
Indels
                                                             Length 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                           43 TTTCAACACTATCGGAATGTGACTGTC 69
                              Zea mays.
Zea mays
                                                                                                                                                                                                                                                                                                             27 TTTTANCNCTATAGGAAAGGGAGGATC 1
                                                                                                                                                                                                                                                                                                                                                                                                                 18;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
                                                                                                   BH643370.1
                                                                                                                                                              1008057C05.1EL_x1 1008 - RescueMu Grid I Zea mays genomic, DNA
                                                                                                                                                                                               BH643370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         submitting the same cDNA sequence redundantly to DBB since 1993. For the abundance information of clones with this sequence in this library and as well as in other 3'-directed libraries, see http://www.imcb.osaka-u.ac.jp/bodymap'. The sequences of the clones represented by this GS sequences is also found there.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1-3,Yamada-oka, Suita, Osaka Pret. 565, Japan
Tel: 06-877-5111(ex.3315)
Bmail: kousaku@imcb.osaka-u.ac.jp
Human Gene Signature, 3'-directed cDNA sequence. We are not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Osaka University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Okubo, K
                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BodyMap; human gene expression database 
Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Organ: blood; Vector: 1-gt-11; Site_1: Eco-RI; Moncoytes were prepared from blood by ficoll-hypaque, percoll and T cell rosetting purification steps (purity: 96 %). mRNA was prepared from activated moncoytes from a patient with rheumatoid arthritis. mRNA was reverse transcribed with MuIV. Using Eco-RI linkers cDNA was transcribed with MuIV. Using Eco-RI linkers cDNA was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               monocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        marked ss-cDNA from activated and non-activated monocevtes. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cloned into 1-gt-11 vector arms. The cDNA library was screened by differential hybidization using radioactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Human adult (K.Okubo)"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                   GI:18672167
                                                                                                                                                                                                                                                                                                                                                                                                                                   14.8%;

 Mismatches

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REFERENCE
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Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Berkeley MGC sequencing project Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                       Unpublished (1999)
Other_ESTs: 2820451.5prime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 TGGGAGCACTCCATTCTGCCTGGGT 29
                                                                       Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                             Mammaila; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 48)
                                                                                                                                                                                                                                                                                                                                                                                            ESI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2820451.3prime NIH_MGC_7 Homo sapiens cDNA clone IMAGE:2820451 3',
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Tel: 650 723 2227
Fax: 650 725 8221
                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reverse complemented post-ligation sequence from source sequence. Plate: 1008057 row: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Walbot, V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: walbot@stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Maize genomic sequences found using engineered RescueMu transposon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Very probable ligation site of ends cut by single endonuclease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            units. For more information on RescueMu, go to the web site www.zmdb.lastate.edu and follow the links for 'RescueMu,' Grid I was grown at Berkeley in 2001 DNA was extracted from leaf punches, double digested using BamHI and BglII, and ligated to form circular plasmids. DH10B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cells were transformed and then screened on LB plates with ampicillin."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site_1: BamHI; Site_2: BgIII; RescueMu is a 4.9 kb, modified maize Mu transposon designed to allow plasmid rescue from total genomic DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mu elements insert preferentially into transcription
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/clone_lib="1008 - RescueMu Grid I"
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72.0%;
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Pred. No. 9.9e+05;
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RESULT 99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       47 GCACTCCAGCCTGGTCGACAGAGTGAGACTCCGTCTCAAAA 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10 GCACTCCATTCTGCCTGGGTGACAATGCCTTGGTTTCAACA 50
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www-bio.llini.gov/bbry/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center Trimming: cross_match from University of Washingtion Genome Center Trimming: cross_match from University of Washingtion Genome Center PHRAP suite. Poly-T identification: patwatch.pl from Berkeley Drosophila Genome Project. University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: 35
contiguous PHRED high quality bases following vector sequence. Very Low Quality Sequence: Trace file contained 48 contiguous distinct peaks following vector sequence. Polyadenylation: Based upon the presence of a XhoI site followed by a run of 14 or more T residues
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Location/Qualifiers
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Roye,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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clone UUGC1M0339C15 F, DNA sequence.
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
                                                                                                                                                                                              University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                               Contact: Robert B.
                                                                                                                                                                                                                                                                                                   Unpublished (2000)
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/clone_lib="NHH_MGC_7"
/tissue_type="small cell carcinoma"
/cell_line="MGC3"
                                                                                                                                                              Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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                                                                                                                                                                                                                                                                          Weiss
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                                                                        IMAGE Consortium (info@image.linl.gov) for further information. Trace considered overall poor quality Insert Length: 816 Std Error: 0.00 Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA905808 49 bp mRNA linear EST 09-JUN-199 oj85e04.sl Soares_NFL_T_GBC_Sl Homo sapiens cDNA clone IMAGE:1505118 3' similar to TR:Q92802 Q92802 HYPOTHETICAL 67.5 KD
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                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROTEIN. ;, mRNA sequence.
                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                         Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     (bases 1 to 49)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (91|473214| jb) AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amploillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
/organism="Homo sapiens"
                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:10090"
/clone="UUGC1M0339C15"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.8; Db 1/,
Pred. No. 1e+06;
"" matches 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EST 09-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Search completed: July 5, 2003, 13:09:21 Job time: 1409 secs
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ORIGIN
                                                                                                                                                                                                                                         Query Match 14.8%; Score 13.8; DB 9; Length 49; Best Local Similarity 63.6%; Pred. No. 1e+06; Matches 21; Conservative 0; Mismatches 12; Indels
                                                                                                                                                             56 GGAATGTGACTGTCGGGCCTCCAGAAAACATTG 88
                                                                                                                   33 GGAATAAACTTGTTGTGTCTCGAAAGAAGATTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                     12 a
                                                                                                                                                                                                                                                                                                                                                                                                       /note="Organ: pooled; Vector: pf773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Ecc RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_GAP_GCBL) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 72909-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref="taxon:9606"
/clone="IMAGE:1505118"
/clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
                                                                                                                                                                                                                                             0; Gaps
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В δ

